

Can Signaling Molecules in Fibroblasts Detect Prodromal Alzheimer's Disease?

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Abstract

Ras/ERK, /p-38, /JNK and PI3K/AKT pathways may mediate amyloid-β-peptide toxicity, neuron death and cognitive decline, modulating oxidative stress and inflammation in AD. Although it is a neurodegenerative disorder, AD affects different systemic molecular mechanisms that are present in patients' peripheral tissues, too. We investigated possible MAPK and PI3K-AKT alterations in fibroblast primary cultures from sporadic AD, MCI and control subjects, testing these molecular pathways as possible AD-related trait markers.

By western blot and phospho-Elisa, we evaluated their epigenetic status and some of their downstream pathways that might be involved in preclinical and early AD stages: the alpha-APP-levels, as a marker of the alpha-secretase activity, the expression of Bax, as an apoptosis and mTOR-autophagy regulator and the phospho-p70S6K status, that modulates protein synthesis, Tau hyper-phosphorylation and mTOR-cell signaling.

While the expression of phospho-ERK1/2 decreased in fibroblasts from mild/moderate AD compared to severe patients' and controls' cells, it did not shown difference in fibroblasts from seven sporadic Parkinson Disease patients compared to HC. Moreover, phospho-ERK1/2 expression and AD severity correlated and including MCI-converter patients to AD cases, this statistical significance was further confirmed. The expression of phospho-AKT also increased in the cells of severe AD patients compared to the fibroblasts of patients with mild/moderate AD, MCI and control subjects and it correlated with the disease severity, too. Phospho-p38- and phospho-JNK-SAP Kinases as well as phospho-p70S6K and Bax increased in cells from AD patients but did not correlate with disease severity, while alpha-APP levels decreased in patients' cells and it only correlated to phospho-ERK1/2 levels in control fibroblasts, indicating the ERK involvement in the physiological metabolism of APP.

Then ERK1/2 and AKT pathways could be suitable to check the disease molecular mechanisms in these peripheral cells from the disease prodromal stage, in particular, ERK-modulation might help to discriminate converter and nonconverter MCI. Furthermore, since fibroblasts show molecular dysfunctions similar to those observed in autopsy brains, these cells might also be useful peripheral targets to investigate and modulate Aβ-related APP metabolism, Tau hyper-phosphorylation and mTOR-cell signaling, preventing the AD progression.

Keywords: ERK and AKT; MMSE; MCI-Converter; Alpha-APP; P70S6Kinase; Tau synthesis; Apoptosis; mTOR-pathway; Personalized drugs

Abbreviations

AD: Alzheimer's Disease; MCI: Mild Cognitive Impairment; HC: Healthy Control; MAPK (JNK, p38 and ERK): Mitogen-Activated Protein Kinase; ERK: Extracellular Signal-Regulated Kinase; JNK: cJun N-terminal Kinase; AKT: Protein Kinase B; PI3K: Phosphatidylinositol-3-Kinase; p-ERK, -JNK,-P38 and -AKT: Phospho-ERK, Phospho-JNK, Phospho-p38 and Phospho-AKT; A β : Amyloid Beta Peptide; ROS: Reactive Oxygen Species; MMSE: Mini-Mental State Examination; APP: Amyloid Precursor Protein; CNS: Central Nervous System

Introduction

During the different phases of Alzheimer's disease (AD), a variety of growth factors and mitogen compounds are modified, mediating their cellular effects through activation of MAPK (JNK, p38 and ERK1/2) and PI3K cascade [1,2]. These pathways regulate many cellular processes that are involved in cognitive functions, too [3-8].

JNK and p38 are activated by a variety of environmental stresses, inflammatory cytokines, growth factors and G-protein-coupled receptors agonists. Furthermore, ERK plays an important role in

J Alzheimers Dis Parkinsonism, an open access journal ISSN: 2161-0460

regulating cell growth and proliferation, and modulating stress and inflammation signals through the activation of p53 and autophagy by mTOR [8,9]. ERK regulates the activities of several transcription factors, such as ELK1 and CREB that play a kay role in neuronal differentiation and long-term memory formation, too.

PI3K-AKT signaling has great importance in the cell cycle process, promoting survival and growth in response to extracellular signals and also fulfills various biological functions including angiogenesis, apoptosis and autophagy, through activation of downstream responses. It also phosphorylates MAP-kinase-kinase-kinases that are the upstream protein regulators of MAPK. Currently, these pathways have been involved in several molecular mechanisms responsible for

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Received April 30, 2020; Accepted June 09, 2020; Published June 16, 2020

Citation: Zoia CP, Compagnoni P, Bazzini C, Ulisse A, Conti E, et al. (2020) Can Signaling Molecules in Fibroblasts Detect Prodromal Alzheimer's Disease? J Alzheimers Dis Parkinsonism 10: 490.

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the AD pathogenic processes, even if the underlying mechanisms and interactions are not yet outlined.

Furthermore, ERK1/2 and AKT may regulate the phosphorylation of p70S6-kinase and its phosphorylation at threonine 389 has been correlated with autophagy modulation, too.

ERK1/2 pathway has also involved in Tau-hyper-phosphorylation, which is evident before the deposition of A β fibrils, through p70S6-kinase activation, a downstream pathway of PI3K/AKT [4,7]. Recently, Li observed harpagoside, a natural anti-inflammatory, improves in rats the A β -induced cognitive impairment via up-regulating BDNF expression, modulating MAPK and PI3K pathways [8,10].

JNK, p38 and ERK1/2 activation has been demonstrated in AD neurons and dystrophic neuritis and in CNS from patients at early disease stages [11-16]. In AD neocortical pyramidal neurons, the neurofilament hyperphosphorylation is accompanied by ERK1/2 activation, that is associated with age-dependent amyloid plaque deposition and increased oxidative stress and the modulation of secretase in AD animal model [17-19]. Pei and colleagues specify that ERK1/2 active forms have been found both in tangle-neurons and in AD neurons that did not show abnormal hyperphosphorylation of Tau [20]. In human CSF, ERK1/2 correlates with total- and phospho-Tau levels and elevated pTau-levels and reduced $A\beta_{1-42}$ concentration are described as molecular biomarkers [21].

ERK1/2 is related to $A\beta$ accumulation, with consequent memory dysfunction in Tg2576 mice and phospho-ERK1/2 is involved in synaptic plasticity and memory formation through CREB-activation [6,22,23]. Pharmacological or molecular inhibition of ERK1/2 and/ or CREB function leads to memory impairment, whereas their genetic overexpression and activation contribute to memory enhancement [5,24]. Rilapladib, an oral inhibitor of lipoprotein associated phospholipaseA2 that can have pro-inflammatory and oxidative properties, is also activated by ERK1/2 phosphorylation and it is currently investigated in phase II trials for AD [25].

In animal model, p38 and JNK activation has also associated with $A\beta$ deposition and tau-hyperphosphorylation and the link between intracellular $A\beta$ and JNK phosphorylation has been confirmed in cortical neurons from sporadic and PS1-linked AD patients[26-28].

Moreover, $A\beta$ toxicity mediates MAPK and PI3K signaling in animal models, neuroblastoma cell line and primary human fibroblast cultures [8,29-31].

In bradykinin-treated fibroblasts from sporadic and familial patients, dysfunctions in ERK1/2 signaling cascade have also detected and correlated to dementia duration [31,32].

Activated ERK1/2 and PI3K/AKT have been shown to directly phosphorylate GSK3 β and GSK3 β -genetic variants are associated to total- and phospho-Tau and A β levels, in AD patients [33,34]. Moreover, fibroblasts are a useful cell model to investigate Tau modifications [35].

To investigate the complex interactions among A β , Tau and MAPK and PI3K/AKT signaling, in patients with different disease severity, we

compared these parameters in primary fibroblasts obtained from mild and moderate/severe patients, Mild Cognitive Impairment (MCI) and Healthy (HC) subjects [36,37]. Although some studies show modulation of these pathways in the autopsy brain of the patients and/or in pathology models, our results indicate a specific ERK and AKT relationship to severity of the disease in patient's fibroblasts and their involvement in A β -dependent mTOR signaling [38]. We suggest that these signaling pathways, playing a key role for the disease, may help to select and target clinical investigations, suggesting some personalized therapeutic strategies [21,39]. Thus, they might help to identify individuals at an early stage or high risk of the disease.

Materials and Methods

Population study

Twenty AD patients (six severe, eight moderate and six mild), 7 MCI subjects and 13 age-related controls took part into the study. Their socio-demographic and clinical features are shown in Table 1. Criteria of the National Institute of Neurological and Communicative Disorders and Stroke and the Alzheimer's Disease and Related Disorders Association (NINCDS-ADRDA) were applied for the diagnosis of probable AD or MCI [40,41]. All patients presented with progressive cognitive impairment predominantly affecting memory and were assessed with brain Computed Tomography (CT) or nuclear magnetic resonance imaging (NMRI), blood tests for excluding secondary dementia (thyroid hormones, folic acid, vitamin B12, VDRL-TPHA and erythrocyte sedimentation rate) and a comprehensive neuropsychological battery. AD patients were considered severe if they had a MMSE score < 12, moderate for a score between 13 and 19 and mild for a score between 20 and 24. Exclusion criteria were psychiatric disorders such as major depression, mania and schizophrenia, a past or present history of substance abuse, neoplastic or haematologic disorders, recent infections or surgery, severe hepatic or renal insufficiency, myocardial infarction or head injury within the last six months and ongoing antiplatelet, anti-inflammatory, anti-neoplastic, corticosteroid or immunosuppressive treatments. No patient had a familial history of dementia. For HC, active medical illness, personal or family history of neurological or psychiatric disorders and alcoholism or drug abuse were ruled out by clinical interview. Absence of cognitive impairment was defined by a MMSE score>26/30. Experiments were undertaken with written consent of each participant and the study was approved by the local Ethics Committee of San Gerardo Hospital, Monza, Italy.

Skin biopsy and primary cell culture

From HC and MCI subjects and patients skin biopsies were performed and cultured into Dulbecco's Modified Eagle's Medium (DMEM, Euroclone, Italy), supplemented with 10% heat inactivated Fetal Calf Serum (FCS), 2 mM glutamine, 100 U/ml penicillin and 100 mg/ml streptomycin. The obtained fibroblasts were plated in dish and they were expanded until they reached the sub-confluence. Then they were collected in cold PBS within 12th trypsin treatment, as previously described [30].

Subjects	Number	Men/Women	Age ± SD	MMSE ± SD
AD	20	9/11	74.2 ± 1.99	16.63 ± 5.09
HC	13	6/7	69.4 ± 1.63	29.22 ± 0.47
MCI	7	4/3	74.6 ± 2.04	27.64 ± 1.03

Table 1: Demographic characteristics and neuropsychological score of tested subjects. APO-E ϵ 4 total distribution in AD (20) and MCI (7) is 11/27. For Age, One-wayanalysis of variance is: p=0.35, for MMSE, AD vs HC p< 0.01 and AD vs MCI p<0.05 and HC vs MCI p>0.05.

To obtain dry pellets, fibroblasts were centrifuged at 250 x g at 4° C for 10 min, then they were preserved at -20°C, until the protein extraction.

Western Blot analysis (WB)

Sub-confluent cell cultures of fibroblasts were collected in cold PBS and lysated in Cell Extraction Buffer, containing protease and phosphatase inhibitors (CEB, Invitrogen, UK) [30]. Proteins of cytosol (25 ug) were separated by electrophoresis on 10-12% SDS polyacrylamide gels and they were blotted on nitrocellulose filter. Blots were blocked 1 h at room temperature on a shaker in 5% no-fat dried milk in TBS-T buffer (50 mM Tris-HCl pH 7.6, 200 mM NaCl, 0.1% Tween 20).

Anti-p38, -JNK, -ERK1/2, -AKT, -p70/S6 and -Bax immunoblot (WB)

Blots were incubated overnight, on a shaker, at 4°C with the primary rabbit antibodies. They were: Anti-phosphorylated p38 [pTpY180/182] and anti-total p38, anti-phosphorylated JNK [pTpY183/185] and anti-total JNK, anti-phosphorylated ERK1/2 [pTpY185/187] and anti-total ERK1/2 (1:500 polyclonal), monoclonal anti-phospho-AKT (1:500), anti-phosphorylated p70 S6-Kinasi [T389] (1:300), anti-phospho-Tau (1:1000, [Ser400/404-Thr403]) and mouse anti-Bax (1:200, Merck-Millipore Italy). All antibodies (Cell Signaling, The Netherlands) were diluted in 5% no-fat dried milk in TBS-T buffer. Mouse anti-actin antibody (1:20000, Merck, Italy) was internal standard.

Peroxidase-linked anti-rabbit/mouse (1:5000/7000 or 1:15000 for actin antibody, Merck, Italy) IgG secondary antibody were incubated for 1 h, at room temperature, on an orbital shaker in TBS-T buffer, containing 3% no-fat dried milk. The presence of signals was detected by chemiluminescent reagents such as ECL Plus Kit (Merck, Italy). The protein signal was capture with a photographic film and imaging densitometer (BioRad, USA) was employed to scan films, which were analyzed by Bio-Rad Multi-Analyst/TM software. The value of signals in WB was expressed as the ratio between the target and actin signals and/or between phosphorylation and total form of the specific kinase, in our experimental condition, no difference was shown in the trend of the two different ratios. Experiments were performed in duplicate or tripled, on different gels.

Total- and phospho-ELISA for MAPK and AKT

In order to detect and quantify the levels of these kinases in protein lysates of fibroblasts, we used the Immunoassay Kits (phospho-ELISA kits, BioSource International, Inc, USA). Total levels of p38, JNK, ERK1/2 and AKT are independent of phosphorylation status, so we used total ELISA kits (BioSource International, Inc, USA) to normalize the phosphorylated p38, JNK, ERK1/2 and AKT amount in the samples.

Cytosol protein extraction was performed in CEB buffer, containing 1 mM PMSF, protease and phosphatase inhibitor cocktail (Merck, Italy, 1:200 and 1:100), for 30 min, on ice. Then lysates were centrifuged at 12000 x g for 10 min at 4° C.

According to the Bio Source assay method, different dilutions of samples were tested for each phosphorylated or total protein detection. The protein absorbance was determined by plate reading at 450 nm (BioRad, USA). The concentrations were calculated comparing the tested absorbance to the specific standard curve values, for each MAPK phosphorylated status and expressed with respect to specific MAPK total status.

Levels of human alpha-APP in cell lysates

In fibroblasts form recruited subjects, APP levels were detected in cytosol protein extracts. By Immunoassay Kit (Merck-Invitrogen, Italy), alpha-APP (α APP) levels were tested. This kit detects soluble α APP which is cleaved by alpha-secretase, but not soluble beta-APP which is cleaved by beta-secretase, as described in commercial data sheet.

Statistical analysis

By Graph Pad Prism program statistical analysis was performed. All results were expressed as mean \pm Standard Deviation (SD). Oneway ANOVA analysis, followed by Bonferroni post-test, was used to determine the significance of differences among the values obtained in the subgroups of recruited subjects. By Pearson's test, possible correlations between protein-kinase expressions and MMSE score or α APP levels were analyzed. By Student-t test, we evaluated the statistical analyses between patients' and HC group.

Results

Stress-activated protein kinase in AD fibroblasts

The phosphorylation state of p38 and JNK was assessed in fibroblasts from AD, MCI and age-matched HC.

By western blot and phospho-ELISA assays, comparable results were shown (Figures 1A and 1B). A significant difference (p=0.03) between patients (1.07 \pm 0.36, ratio value in WB) and HC (0.47 \pm 0.23, ratio value in WB) was observed, indicating a 100% increase of p38-phosphorylation in AD subjects, from the earliest stages of the disease. A trend to p-p38 increase was shown in MCI subjects (0.66 \pm 0.20, ratio value in WB, p=0.21 vs HC and p=0.06 vs AD), too. By WB, anti-phospho-JNK antibody displayed specific and 100% increased immunoreactivity in AD (1.45 \pm 0.32, ratio value in WB) compared to control fibroblasts (0.78 \pm 0.28, p<0.05) and in MCI fibroblasts (1.20 \pm 0.42, p>0.05) (Figure 1C). By phospho-ELISA, comparable results were shown (Figure 1D). There is no correlation between p-JNK or p-p38 and disease severity.

The RAS-ERK and PI3K-mTOR pathways and disease severity in AD fibroblasts

By WB, the examples of p-ERK modulation were shown in fibroblasts from MCI and AD compared to control fibroblasts (Figure 2A). By phospho-ELISA assay, ERK1/2 phosphorylation reduced by approximately 50% in patients, precisely in fibroblasts from mild and moderate AD (0.021 \pm 0.006 p/tot, p<0.001) compared to HC cells $(0.050 \pm 0.020 \text{ p/tot})$. Furthermore, there was a statistically significant increase of p-ERK1/2 in cells from severe AD patients (0.053 ± 0.012 p/tot) compared to fibroblasts from mild and moderate patients (p<0.01). No difference was observed between severe AD and HC, while p-ERK1/2 decreased in MCI fibroblasts, too (0.026 \pm 0.017 p/tot, p <0.05 vs HC) (Figure 2B). We also observed a significant inverse correlation between p-ERK1/2 and MMSE score (r= -0.596, p=0.01), indicating a direct correlation with disease severity. Including the p-ERK values and corresponded MMSE score of four converters MCI, the correlation was maintained (Figure 2C). Moreover, in non-converter-MCI subjects, no correlation between p-ERK levels and MMSE score was shown.

As well as ERK1/2, AKT kinase is involved in cell survival and it modulates mTOR-p70S6K pathway. In patients' fibroblasts, p-AKT increased by 40% or 50% in western blot (p=0.02) or ELISA (p=0.0008),

J Alzheimers Dis Parkinsonism, an open access journal ISSN: 2161-0460





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respectively vs HC cells (Figures 3A-B). In MCI, p-AKT status was reduced compared to severe AD (p<0.001) and it was decreased with respect to mild/moderate patients (p<0.01), too. In addition, there is a direct correlation between phospho-AKT expression and disease severity, evaluated by MMSE score (r=-0.63, p=0.03) (Figure 3C).

Regulation of Tau metabolism and apoptosis in patients' fibroblasts

We also tested the phosphorylation of p7086-kinase that is regulated by ERK1/2 and AKT pathway and its phosphorylation at threonine 389 has been correlated with autophagy inhibition, too. In particular, the 70kDa isoform regulates the activation of Tauprotein synthesis and its phosphorylation, as described in CNS cells. While the 85KDa nuclear isoform modulates the transcriptional mechanisms.

In fibroblasts, our results showed an increasing trend of phosphop70S6K in AD patients (p<0.001 vs HC) and it is interesting to note that the predominant isoform is the cytosolic one (70KDa) (Figure 4A). Our results also indicated the increase of pTau- Ser400/Thr403/Ser404 in fibroblasts from AD patients compared to the cells of HC (p<0.05) (dashed box) (Figure 4).

Moreover, in a subgroup of patients and HC, it was detected an increase (p<0.05) of the cytoplasmic levels of Bax (Figure 4B). This result indicates the triggering of the apoptotic mitochondrial pathway [42]. Bax is also a downstream pathway that both ERK1/2 and AKT regulate, in fact, it modulates apoptosis and autophagy signaling.

ERK and APP metabolism

Since the ERK1/2 signaling might be linked to APP metabolism, we also tested the cytosolic α APP levels in fibroblasts from recruited subjects [43]. A 60% reduction of the α APP levels were observed in AD patients, in particular in fibroblasts from severe patients compared to HC (p=0.003) (Figure 5A). In MCI fibroblasts, α APP distribution was heterogeneous. Moreover, phospho-ERK1/2 and α APP correlation was very significant in HC (r=0.915), while in MCI subjects and AD patients this correlation was lost (Figure 5B).

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fibroblasts from AD patients (0.05 \pm 0.01) and HC (0.0017 \pm 0.0005), *p<0.005 vs HC. In the graphic, there is the ratio of their O.D. value.







Discussion

Various research groups have turned their attention to the signal transduction pathways involved in the pathway of GSK-3 β , such as MAP kinase, PI3K/AKT and PKC. They play a role in mediating the toxic effect of A β in the regulation of Tau hyper-phosphorylation, this toxicity may also modulate the mechanisms of synaptic and neuronal degeneration [34,44-48].

It knows that several systemic mechanisms and pathways may be investigated in fibroblasts, reflecting the biochemical and molecular alterations that have been observed in CNS [30-33,35]. Since in the brain of AD patients, MAPK and PI-GSK3/AKT modulations seem related to the alterations and deposition of A β and Tau, we investigated these specific pathways in primary cultures from human fibroblasts. In this way, we considered systemic but specific molecular mechanisms involved in AD, particularly in the early phase of disease.

Since fibroblasts are independent of the postmortem changes, they can be suitable for pharmacological and molecular studies, too [49]. Moreover, in these cells the pathogenic mechanisms, involved in AD progression, may be investigated as possible therapeutic targets [32]. Following autopsy, the same PKC alterations have been described in both fibroblasts and brain, therefore it thinks these kinase-linked molecular changes and/or their epigenetic modifications may also be maintained in these peripheral cells [31].

In this work, the data show affected MAPK and AKT pathways, in fibroblasts from AD patients and MCI subjects compared to HC, highlighting the role of these pathways in A β induced-oxidative stress, APP metabolism, Tau phosphorylation and their involvement in the etiopathology and progression of disease, modulating inflammation and mTOR-signaling.

In particular, since the early stages of AD, the stress activated phosphokinases p38 and JNK increased their phosphorylation and a clear upward trend has also shown in MCI. These modifications may be a response to cell bio-energetic impairment but they did not correlate to the disease severity. Moreover, the involvement of these stressactivated kinases, in the early stage of disease, might also be responsible for the reduced Energy-Dependent Activity of Glutamate Transporters (EAATs), justifying in part the decreased glutamate uptake, previously observed in fibroblasts from AD patients, in the absence of EAATs modifications both to mRNA and protein expression [39,43].

In addition, the signal transduction pathway mediated by p38 is involved in the alteration of synaptic plasticity caused by

 $A\beta$, suggesting a role of this kinase at CNS level, in the decline of the mnemonic functions, features of the disease [39]. Bodles and colleagues hypothesized that activated microglia, observed in the early stage of the disease, is able to phosphorylate JNK and p38, inducing the production of factors that are pro-inflammatory and toxic to neurons [50]. Moreover, these kinases are jointly responsible for Tau phosphorylation that we observed to be dysregulated in AD fibroblasts, too [51].

Different authors also report ERK1/2 activation in AD neurons and dystrophic neuritis [13-15]. Khan described ERK1/2 as a putative peripheral biomarker for AD, in response to the inflammatory signal bradykinin [31]. Furthermore, p-ERK1/2 level increased in CSF from AD patients, as described by Spitzer and colleges [21].

In line with these papers, we also underlined the specific ERK1/2 involvement in AD fibroblasts. In detail, we observed that ERK1/2 phosphorylation depends on the disease severity: ERK1/2 was downphosphorylated in MCI, mild and moderate AD patients with respect to healthy subjects, while p-ERK1/2 levels increased in severe AD cells and they were higher than HC levels. Surprisingly, the inverse correlation between MMSE score and ERK1/2 phosphorylation strengthened the modulation of ERK1/2 in different disease stages. In fact, over at least three years, the four MCI subjects that converted to AD patients have kept this correlation, while it did not show in non-converter-MCI subjects (Figure 2C). For the first time, this correlation has been observed in fibroblasts and we propose this parameter might help to predict the evolution of MCI status. Integrating with neuroimaging investigations, a specific longitudinal study could confirm this hypothesis, suggesting p-ERK as an economical and useful peripheral marker. In fact, we tested p-ERK1/2 levels in fibroblasts from patients with Parkinson's disease, but no difference showed (p>0.05, n=7 data not shown).

ERK1/2 also modulates inflammatory stimuli thus its phosphorylation, in the later stages of the disease, could account for the involvement of inflammatory mechanisms in AD pathogenesis [31]. This phenomenon could also suggest that a compensatory mechanism may occur in severe AD fibroblasts, resulting in the cell attempt to balance the oxidative stress that A β induces from the early stages of disease. On the other hand, an unbalance between kinases and phosphatases system may occur in fibroblasts from severe AD patients. In fact, in fibroblasts from severe patients, Zhao and colleagues demonstrated decreased levels of phosphatase 2A (PP2A), inducing a prolonged ERK1/2 activation [33]. This protein phosphatase and PP1

phosphatase also decrease in the brains of AD patients and they regulate Tau phosphorylation [52]. Moreover, the up regulation of MKP-1, that is an essential negative regulator of MAPKs by dephosphorylating them at both tyrosine and threonine residues, reduces A β generation and alleviates cognitive impairments in Alzheimer's disease models [53].

Owing to the decreased phosphatase activity in the AD brain, the dephosphorylation of these kinases and Tau would be compromised, allowing Tau to be abnormally hyper phosphorylated at multiple sites. In fact, Tau is phosphorylated at more than 21 sites in AD brain and ERK1/2 has been referred to be responsible for Tau phosphorylation, too [7,26]. The p70 S6-kinase is a downstream protein of ERK1/2 pathway and its cytosolic isoform (70kDa) is involved in the activation of Tau-protein synthesis and its phosphorylation, as described in CNS [54]. Since in AD fibroblasts, we observed a specific increased in 70KDa isoform, we hypothesized an involvement in Tau post-translational regulation for ERK1/2-activated p70S6K. We have tested this hypothesis and our data indicate Tau hyper phosphorylation in AD cells.

The data obtained in our laboratory shown that the pathway regulated by PI3K/AKT is up-regulated in fibroblasts from severe AD patients compared to HC cells, in the early stages of the disease, a trend towards increased p-AKT has also observed. An opposite trend has been showed in MCI. Moreover, a significant correlation between AKT phosphorylation and the disease severity was observed, too. This finding could also be explained by the involvement of AKT in GSK-3 β phosphorylation, the key kinase involved in Tau hyper-phosphorylation [55]. Several authors have demonstrated AKT activation in brains from AD and our results are in agreement with these data [33].

The activation of PI3K/AKT signaling pathway would lead to phosphorylation of AKT cascade, i.e. p70S6K, Tau, mTOR, Bax, modulating A β -oxidative damage, protein expression and their epigenetic regulation, autophagy and apoptosis pathways. The phosphorylation of p70S6K at threonine 389 has been used as a hallmark of activation by mTOR and correlated with autophagy inhibition in several diseases [34,36-38]. Also in fibroblasts, the increase of p-AKT and p-p70S6K could support the involvement of this signaling pathway in AD pathogenesis, mirroring CNS results.

Moreover, the induction in the cytoplasmic levels of Bax highlights the triggering of apoptotic mitochondrial pathway [43]. In the early stages of the disease, this result could be due to the activation of p38 and JNK (SAP Kinase) that integrate with the modulation of ERK1/2 and AKT signal transduction pathways (Figure 6). The assessment of p-AKT and p-ERK1/2 levels in fibroblasts from patients may be a molecular support to the clinical investigations. It might also be useful to studyi the relationship between the modulation of AKT and ERK1/2 phosphorylation and Tau protein expression [56].

The secretion of α APP by a-secretase is a highly regulated second messenger pathway that involves several protein kinases including MAPK, PKC, PKA, etc. [44,45]. Here we showed a decreased α APP levels in fibroblasts from AD and some MCI, compared to HC.

Furthermore, the correlation between α APP detection and ERK1/2 phosphorylation in HC fibroblasts evidences a straight ERK1/2 involvement in α APP metabolism, that is lost during the disease progression, as shown in AD brain [43]. No correlation was observed between SAPKs or AKT and α APP levels.

Instead, AKT seems assumes a more specific involvement in Tau phosphorylation, autophagy and apoptosis modulation.

activity and reduced αAPP levels in fibroblasts from patients, right from the earliest disease phases, while in severe AD a different trend appears. All these evidences in our ex-vivo cellular model show a key role for ERK1/2 and AKT, in the disease physiopathology, supporting and clarifying some indications obtained from the brain autopsy of AD patients [57,59].
 Conclusion

 Our studies support the validity of fibroblasts as peripheral ex vivo model for studying the pathogenesis of AD. They showed the kay involvement of p-ERK1/2 and p-AKT in relation to disease severity and, for the first time, we observed that the phosphorylation status of ERK

might differentiate between MCI-converter and MCI non-converter. Moreover, our results underline that the regulation of mTOR signaling in AD also has a kay role in these peripheral cells, facilitating the approach of biomolecular and pharmacological research.

On the other hand, recent data have showed the physiologic

aAPP function in limiting neuron damage and death, in response to

neurotoxic stress conditions: aAPP is able to stimulate AKT activity

in wild-type fibroblasts, human neuroblastoma cells and neurons

[43,56,57]. According to these data, we observed a decreased AKT

Nowadays, skin fibroblasts are the most suitable source of induced Pluripotent Stem (iPS) cells, too [58,60]. These cells and the signaling pathways, that we have investigated, are considered for the development of cell models for some neurodegenerative diseases and drug screening. From this point of view, our study might be a pioneering approach to investigate any mechanisms related to Alzheimer's disease and it might also be useful to test possible personalized therapy.

Declarations

Potential conflicts of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest. The research has been given ethical approval in accordance with the Declaration of Helsinki.

Acknowledgement

This work was partially supported by:

The Italian Ministry of University and Research [grant n°CTN01_00177_165430 ALISEI-IVASCOMAR, 2013-2017: Cluster project for Identification, Validation and commercial development of new diagnostic and prognostic biomarkers for complex diseases],

The European Community grant agreement n°212043: NAD, 7°Framework Programme, 2007-2013.

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