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In A β PP-overexpressing cultured human muscle fibers proteasome inhibition enhances phosphorylation of A β PP751 and GSK3 β activation; effects mitigated by lithium and apparently relevant to sporadic inclusion-body myositis

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Abstract

Muscle fiber degeneration in sporadic inclusion-body myositis (s-IBM) is characterized by accumulation of multiprotein aggregates, including aggregated amyloid- β -precursor protein 751 (A β PP⁷⁵¹), amyloid- β (A β), phosphorylated tau (p-tau), and other “Alzheimer-characteristic” proteins. Proteasome inhibition is an important component of the s-IBM pathogenesis. In brains of Alzheimer disease (AD) patients and AD transgenic mouse models, phosphorylation of neuronal A β PP⁶⁹⁵ (p-A β PP) on Threonine⁶⁶⁸ (T⁶⁶⁸) (equivalent to T⁷²⁴ of A β PP⁷⁵¹) is considered detrimental because it increases generation of cytotoxic A β and induces tau phosphorylation. Activated glycogen synthase kinase3 β (GSK3 β) is involved in phosphorylation of both A β PP and tau. Lithium, an inhibitor of GSK3 β , was reported to reduce levels of both the total A β PP and p-A β PP in AD animal models. In relation to s-IBM, we now show for the first time that: 1. In A β PP-overexpressing cultured human muscle fibers (human muscle culture IBM model: a) proteasome inhibition significantly increases GSK3 β activity and A β PP phosphorylation; b) treatment with lithium decreases i) phosphorylated-A β PP; ii) total amount of A β PP, iii) A β oligomers, and iv) GSK3 β activity; and c) lithium improves proteasome function. 2. In biopsied s-IBM muscle fibers, GSK3 β is significantly activated and A β PP is phosphorylated on Thr⁷²⁴. Accordingly, treatment with lithium, or other GSK3 β inhibitors, might benefit s-IBM patients.

Keywords

Inclusion-body-myositis; phosphorylated amyloid- β precursor protein; GSK3 β ; cultured human muscle fibers; lithium chloride; proteasome

INTRODUCTION

Sporadic-inclusion body myositis (s-IBM) is the most common progressive muscle disease associated with aging. Its progressive course gradually leads to pronounced muscle weakness and wasting, resulting in severe disability (Askanas and Engel 2008). The exact pathogenesis of s-IBM is not known, and there is no enduring treatment.

An intriguing aspect of s-IBM is that its muscle-fiber phenotype shares several molecular abnormalities with Alzheimer disease (AD) brain, including accumulation of amyloid- β

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precursor protein (A β PP) and of its cytotoxic amyloid- β 42 (A β 42) fragment (recently reviewed in Askanas and Engel 2008; Askanas *et al.* 2009). Both s-IBM and AD affect aging individuals, but they are clinically distinct.

Hallmarks of s-IBM muscle fibers are: 1) vacuolated muscle fibers having intra-myofiber accumulations -- mainly in non-vacuolated cytoplasm -- of congophilic, ubiquitinated, multi-protein aggregates that contain, in addition to A β , phosphorylated tau in the form of paired helical filaments, and several other “AD characteristic” proteins; and 2) mononuclear-cell inflammation (Askanas and Engel 2001; Askanas and Engel 2008; Dalakas 2008). Currently there is no consensus regarding whether the degenerative or inflammatory component plays a more significant role in s-IBM pathogenesis. Against a primary role of inflammation is the wide recognition that all types of immunosuppressive treatment fail to reverse, or enduringly halt, disease progression of s-IBM (in contrast to polymyositis). And, there is increasing agreement that A β 42 is an important pathogenic factor leading to muscle-fiber destruction (Askanas and Engel 2007; Askanas and Engel 2008; Dalakas 2008). This proposed mechanism has been supported by models involving cultured human muscle fibers and transgenic mice. For example, long-term overexpression of wild-type full-length 751A β PP in cultured normal human muscle fibers produced in them several aspects of the s-IBM cellular phenotype, including vacuolization, congophilic amyloid inclusions, cytoplasmic 6–10 nm amyloid-like filaments, nuclear paired helical filaments, mitochondrial cytochrome oxidase deficiency and mitochondrial morphological abnormalities (Askanas *et al.* 1996; Askanas *et al.* 1997). Transgenic mouse models based on overexpressing A β PP or its fragment in skeletal muscle produced some aspects of the IBM pathology (Fukuchi *et al.* 1998; Jin *et al.*, 1998; Kitazawa *et al.* 2006). In s-IBM muscle fibers, there is increased production (Sarkozi *et al.* 1993; Guerin *et al.* 2007) and accumulation of aggregated amyloid- β precursor protein-751 (A β PP⁷⁵¹) (Askanas *et al.* 1993), and preferential accumulation of the cytotoxic A β 42 form (Vattemi *et al.* 2009). Accordingly, we postulate that methods reducing muscle-fiber A β accumulation may benefit s-IBM patients. Furthermore, probably-pathogenic oxidative and ER stresses, and proteasome inhibition, also have been demonstrated in s-IBM muscle fibers (Vattemi *et al.* 2004; Fratta *et al.* 2005; Nogalska *et al.* 2006; Terracciano *et al.* 2008).

In AD brain, increased phosphorylation of neuronal A β PP⁶⁹⁵ on Threonine⁶⁶⁸ has been demonstrated (Lee *et al.* 2003; Shin *et al.* 2007), and in brains of AD mouse models, it was considered to be detrimental by increasing generation of A β and inducing tau phosphorylation (Shin *et al.* 2007). Active glycogen synthase kinase 3 β (GSK3 β) was proposed to have an important role in AD pathogenesis because it was shown to modulate phosphorylation of both tau (Anderton *et al.* 2001) and of A β PP on Threonine⁶⁶⁸ (Aplin *et al.* 1996).

Lithium is a reversible inhibitor of GSK3 β (Jope 2003, Klein and Melton 1996; Avila and Hernandez 2007). It presumably acts either a) directly, competing with magnesium in forming a complex with ATP, or b) indirectly, by increasing phosphorylation of GSK on Serine 9, which leads to GSK3 β inactivation (Jope 2003). In AD and IBM mouse models, inhibition of GSK3 β by lithium chloride decreased phosphorylated A β PP and p-tau (Rockenstein *et al.* 2007; Kitazawa *et al.* 2008), respectively.

To explore molecular pathogenic mechanisms in s-IBM, we experimentally modified the cellular micro-environment of cultured human muscle fibers (CHMFs) to mimic various aspects of the s-IBM pathogenesis, thereby producing experimental IBM-Cultured-Human-Muscle-Fiber Models. These models have proved very useful in our previous studies (Askanas *et al.* 1996; Askanas *et al.* 1997; Fratta *et al.* 2005; Nogalska *et al.* 2006, Nogalska *et al.* in press; Wojcik *et al.* 2006).

In the present study we asked whether: 1) in s-IBM muscle fibers, A β PP is phosphorylated and GSK3 β is activated; and 2) in the established s-IBM model involving overexpression of A β PP in cultured human muscle fibers a) imposed proteasome inhibition or experimental induction of ER stress influences A β PP phosphorylation and GSK3 β activity; b) treatment with lithium decreases i) GSK3 β activity, ii) total and phosphorylated A β PP, and iii) A β oligomers.

EXPERIMENTAL PROCEDURES

Muscle Biopsies

Studies were performed on fresh-frozen diagnostic muscle biopsies obtained (with informed consent) from 12 s-IBM, 2 polymyositis, 1 morphologically-nonspecific myopathy, 2 ALS, 2 peripheral neuropathy, and 12 normal controls age-matched to the s-IBM patients (normal muscle biopsies were of patients who, after all tests were performed, were considered free of muscle disease). s-IBM patients were ages 55–79 years, median age 67; normal control patients were ages 53–84, median age 71. Diagnoses were based on clinical and laboratory investigations, including our routinely-performed 16-reaction diagnostic histochemistry of muscle biopsies.

All s-IBM biopsies met s-IBM diagnostic criteria, as described (Askanas and Engel 2001). Not all studies were performed on all biopsies (details below).

Light-Microscopic Immunocytochemistry

Immunofluorescence was performed as described (Askanas *et al.* 1993; Vattemi *et al.* 2004; Fratta *et al.* 2005; Nogalska *et al.* 2006) on 10- μ m-thick transverse sections of 5 s-IBM, 2 age-matched normal, and 7 disease-control muscle biopsies (specified above), using a well-characterized polyclonal antibody specifically recognizing the A β PP⁷⁵¹ isoform only when phosphorylated on Threonine at position 724 (p-A β PP^{T724}), equivalent to Threonine⁶⁶⁸ of the A β PP⁶⁹⁵ isoform (Cell Signaling, Danvers, MA), diluted 1:50. This antibody was produced by immunizing rabbits with a synthetic phosphopeptide corresponding to the residues surrounding Threonine 688 of human A β PP⁶⁹⁵ (corresponding to 724 position on isoform 751) (Ando *et al.* 1999; Muresan and Muresan 2004). To block non-specific binding of an antibody to Fc receptors, sections were pre-incubated with normal goat serum diluted 1:10, as described (Askanas *et al.* 1993; Nogalska *et al.* 2006). Controls for staining specificity were (i) omission of the primary antibody, or (ii) its replacement with non-immune sera or irrelevant antibody. These were always negative.

Combined Immunoprecipitation/Immunoblot Procedure in s-IBM biopsies

To confirm the specificity of the immunocytochemical reaction in s-IBM muscle fibers, a combined immunoprecipitation/immunoblot technique was performed in three s-IBM biopsies, as detailed previously (Vattemi *et al.* 2004; Fratta *et al.* 2005). In brief, 100 μ g of total muscle protein were immunoprecipitated in precipitation-buffer containing 10 μ g of 6E10 antibody (Covance, Princeton, NJ), which on immunoblots recognizes total A β PP and A β . The immunoprecipitated complex, containing IgG antibody along with its bound target antigen and all proteins bound to that antigen, was pulled down using Protein G Sepharose 4 Fast Flow (Amersham) during 4 h of incubation at 4°C. The solution was then centrifuged for 5 min (16,000 \times g at 4°C) and the supernatant removed. The precipitated sepharose immuno-complexes were washed three times with the precipitation buffer by centrifuging 5 min each (16,000 \times g at 4°C). Immunoprecipitates were then electrophoresed and immunoprobed with the rabbit polyclonal anti-p-A β PP^{T724} antibody, followed by an appropriate secondary antibody, and developed using the Western Breeze anti-rabbit chemiluminescence kit (Invitrogen, Carlsbad, CA). To confirm specificity of the physical

association identified by the immunoprecipitation-immunoblot reaction, primary antibodies were omitted from the immunoprecipitation solution. (Control biopsies were not used for the immunoprecipitation experiments because they do not contain enough A β PP to be demonstrated by this technique.)

Immunoblotting

To evaluate whether GSK3 β is activated in s-IBM patients, muscle homogenates of 6 s-IBM and 6 age-matched control biopsies were immunoblotted, as recently detailed (Vattemi *et al.* 2004; Nogalska *et al.* 2006; Nogalska *et al.* 2007; Terracciano *et al.* 2008). In brief, 20 μ g of protein were loaded into 4–12% NuPAGE gels (Invitrogen) and electrophoretically separated. After electrophoresis, samples were transferred to a nitrocellulose membrane. To prevent non-specific binding of the antibodies, the nitrocellulose membranes were blocked in Blocking Reagent (Invitrogen). They were then incubated overnight at 4 °C with a primary antibody. Antibodies used were against a) total GSK3 β (Santa Cruz Biotechnology, Santa Cruz, Ca), diluted 1:50, b) active-GSK3 β phosphorylated at Tyrosine 216 (Novus Biologicals, Littleton, CO), diluted 1:250, and c) inactive-GSK3 β phosphorylated at Serine 9 (Cell Signaling), diluted 1:250. Blots were developed using the Western Breeze chemiluminescent kit (Invitrogen), or ECL Western Blotting Analysis System (Amersham Biosciences, Piscataway, NJ) in combination with horseradish peroxidase (HRP)-conjugated secondary antibody (Jackson ImmunoResearch Laboratories, West Grove, PA). Protein loading was evaluated by the actin band; the results were related to both active and inactive GSK3 β and expressed per total GSK3 β .

Cultured Human Muscle Fibers (CHMFs) and overexpression of A β PP-gene

Primary cultures of normal human muscle were established, as routinely performed in our laboratory (Askanas and Engel 1992), from archived satellite cells of portions of diagnostic muscle biopsies of patients who, after all tests were performed, were considered free of muscle disease. We established 15 culture sets, each from satellite cells derived from a different muscle biopsy. All experimental and control conditions were studied on sister cultures in the same culture set. Not all studies were performed on every set. Twenty days after myoblast fusion was completed, a 3 Kb 751 A β PP-cDNA was transferred into the well-differentiated myotubes using a replication-deficient adenovirus vector at 0.3×10^8 pfu/ml culture medium, as detailed (Askanas *et al.* 1996; Askanas *et al.* 1997; Wojcik *et al.* 2006).

Inhibition of proteasome or ER-stress induction in cultured human muscle fibers overexpressing A β PP

Three days after A β PP gene-transfer, CHMFs were treated with 1 μ M epoxomicin (Biomol Research Laboratories, Plymouth Meeting, PA), an irreversible proteasome inhibitor (Meng *et al.* 1999), for 24h. For comparison, other A β PP-overexpressing cultures were treated for 24 h with ER-stress inducers, either a) tunicamycin, which inhibits N-glycosylation, 4 μ g/ml (Sigma Co, St Louis, MO), or b) thapsigargin, 300 nM (Sigma Co), which inhibits ER calcium-ATPase (Lee, 2005). The doses used were as in our previous experiments (Fratta *et al.* 2005; Nogalska *et al.* 2007; Nogalska *et al.* in press).

Lithium chloride treatment of experimental cultures

Because lithium chloride (LiCl) is known inhibitor of GSK3 β (Jope 2003, Klein and Melton 1996; Avila and Hernandez, 2007), A β PP-overexpressing cultures in 12 independent experiments were treated with 5mM LiCl (Sigma Co) for 48 h. Because human muscle material from which those primary cultures are derived is sparse, we were not able to evaluate wide ranges of lithium dosage. In the present study, we chose a 5mM dose, since that dose was shown effective by others (Kim and Thayer 2009; Yeste *et al.* 2007). In our

preliminary experiments, we evaluated doses of 1, 5, 10, 15 and 20 mM, and found that 5mM was effective without causing any obvious adverse reactions (the 1mM dose was not effective).

Immunoblotting of CHMFs

After treatments (detailed above and below), experimental cultures and their sister-control cultures were harvested and processed for immunoblotting, as detailed previously (Nogalska *et al.* 2007; Wojcik *et al.* 2006). We evaluated total A β PP, pA β PP^{T724}, and GSK3 β with the antibodies described above. 6E10 antibody was used in the concentrations 1:1000 for evaluating total A β PP, and 1:300 to visualize A β oligomers. Protein-loading was evaluated by the actin band. Quantification of immunoreactivity was performed by densitometric analysis using NIH Image J-1.310 software.

Measurement of Proteasome Activity

Proteasome activity was measured as described (Fratta *et al.* 2005). Three main proteasome activities were determined by evaluating the cleavage of specific fluorogenic substrates (Shringarpure *et al.* 2003). 5 LiCl-treated and 5 non-treated A β PP-overexpressing culture sets were homogenized in 20 mmol/L Tris-HCl, pH 7.2, containing 0.1 mmol/L EDTA and 1 mM of fresh dithiothreitol (Meng *et al.* 1999, Demasi *et al.* 2001), centrifuged, the supernatant collected, and protein concentration determined using the Bradford method. Subsequently, 45 μ g of protein from cultured muscle were incubated in 100 μ mol/L fluorogenic substrates for the three different protease activities: Z-Leu-Leu-Glu-AMC (substrate II) for peptidyl glutamyl-peptide hydrolytic (PGPH) activity (Biomol, Plymouth Meeting, PA); Suc-Leu-Leu-Val-Tyr-AMC (substrate III) for chymotrypsin-like (CTL) activity (Calbiochem, Gibbstown, NJ); and Z-Ala-Arg-Arg-AMC (substrate VI) for trypsin-like activity (Calbiochem). Fluorescence emission was excited at 355 nm and recorded at 405 nm. Proteasome activities were expressed per 20s β 2 proteasome subunit performed by immunoblotting each culture set using an anti-20s β 2 mouse monoclonal antibody diluted 1:500 (Biomol).

Statistical Analysis

In all experiments these analyses were performed using Student's t-test. Significance level was set at $p < 0.05$. For all groups data are reported as means \pm SEM.

RESULTS

In s-IBM, muscle fibers A β PP was phosphorylated on Threonine 724

By immunocytochemistry, pA β PP^{T724} was accumulated in the form of various-sized aggregates located mainly in the non-vacuolated regions of cytoplasm, of approximately 80% of s-IBM-vacuolated muscle fibers in all patients (Fig. 1A). None of the normal or disease-control muscle biopsies contained pA β PP^{T724}-immunoreactivity. Endomysial and perivascular mononuclear-cells present in s-IBM and polymyositis muscle biopsies did not contain pA β PP^{T724} immunoreactivity. Muscle blood vessels did not contain pA β PP^{T724} immunoreactivity.

To confirm the specificity of the immunocytochemical staining in s-IBM, we immunoprecipitated three s-IBM muscle-biopsy homogenates with antibody 6E10 (which on immunoblots recognizes total A β PP as well as A β oligomers), and we immunoprobed the blots with the specific anti-pA β PP^{T724} antibody. Immunoprecipitation/immunoblotting confirmed the existence of A β PP phosphorylated on Threonine⁷²⁴ in s-IBM muscle (Fig. 1B).

Active form of GSK3 β was increased in s-IBM muscle fibers—GSK3 β activity is regulated mainly by phosphorylation on Tyrosine 216 (Y²¹⁶) which increases its activity, and on Serine 9 (Ser⁹) which inhibits its activity (Bhat *et al.* 2000; Lochhead *et al.* 2006).

In 6 s-IBM and 6 age-matched normal-control muscle biopsies, we evaluated GSK3 β on immunoblots, using antibodies specifically recognizing total GSK3 β , p-GSK3 β ^{Y216}, and GSK3 β ^{Ser9} (studied in duplicate samples) after their normalization to actin. In s-IBM muscle biopsies, total GSK3 β was not changed as compared to normal controls, whereas the amount of active p-GSK3 β ^{Y216} expressed per total GSK3 β was increased 1.5 fold ($p < 0.05$) (Fig 2). (Inactive p-GSK3 β ^{Ser9} was decreased 1.6 fold in s-IBM muscle biopsies, but this did not reach p-value significance [data not shown].)

Proteasome inhibition increased p-A β PP^{T724} and induced activation of GSK3 β in our A β PP-overexpressing cultures (human-muscle-culture IBM model)

To explore possible mechanisms involved in A β PP phosphorylation and GSK3 β activation in s-IBM muscle fibers, we utilized our experimental model consisting of primary cultures of normal human muscle fibers, with experimental modifications of the intracellular micro-environment to mimic aspects of the s-IBM muscle-fiber milieu. These manipulations involved: overexpressing the A β PP-gene, inhibiting the 26S proteasome, and inducing endoplasmic reticulum (ER) stress.

In 12 independent experiments, immunoblots of cultured human muscle-fiber homogenates, using the phospho-specific antibody recognizing p-A β PP^{T724}, showed that: a) exposure of the A β PP-overexpressing (A β PP⁺) cultured human muscle fibers to the proteasome inhibitor epoxomicin caused 6.5 fold ($p < 0.005$) increase of p-A β PP^{T724}, while b) ER stress inducers tunicamycin and thapsigargin had no effect (Fig.3). (Proteasome-inhibition increased total A β PP 3 fold (< 0.01) [not shown]). As in our previous studies, A β PP from our A β PP⁺-Culture IBM model muscle-fibers migrates as a “mature” 130kDa band and an “immature” 115 kDa band.

To determine whether the proteasome-inhibition-induced increase of p-A β PP^{T724} is associated with GSK3 β activation, we performed immunoblots on 6 independent tissue-culture experiments using the same 3 anti-GSK3 β antibodies as described above for human muscle biopsies. We found that after proteasome-inhibition active p-GSK3 β ^{Y216} was increased 1.7 fold ($p < 0.05$) (Fig. 4), while the total GSK3 β and GSK3 β ^{Ser9} were not affected (data not shown). Since the ER stress did not influence A β PP phosphorylation, we did not evaluate its influence on GSK3 β activation.

Treatment with Lithium inhibited GSK3 β , and decreased total A β PP, p-A β PP^{T724} and A β -oligomers in our culture IBM model muscle-fibers—Lithium is known to inhibit GSK3 β by increasing its inactive p-GSK3 β ^{Ser9} form. When in 12 independent experiments we treated A β PP-overexpressing cultured human muscle-fibers with 5mM LiCl, the inactive p-GSK3 β ^{Ser9} form of GSK3 β was increased 2.2 fold ($p < 0.01$) (Fig.5). That lithium treatment decreased total A β PP 30% ($p < 0.005$), p-A β PP^{T724} 50% ($p < 0.0001$), and A β oligomers (8–25kDa) 25% ($p < 0.05$) (Fig.6). Because in different experiments LiCl decreased A β oligomers of different molecular weights, we calculated all oligomers together.

Lithium induced proteasome activity—In view of the results obtained in our IBM-culture model indicating that proteasome inhibition increased both p-A β PP^{T724} and total A β PP, we asked whether our observed decrease of them by lithium might be, at least partially, influenced by a lithium enhancement of proteasome activity. Therefore, we studied three main proteasome activities in 5 independent sets of the IBM-culture models. We found

that in lithium-treated cultures, all three proteasome enzymatic activities were increased; PGPH activity 2 fold ($p<0.005$), CTL activity 1.3 fold ($p<0.05$), and TL activity 1.2 fold ($p<0.05$), as compared to the sister-non-treated controls, (Fig.7).

DISCUSSION

While A β PP⁶⁹⁵ is specifically present in neuronal cells (da Cruz e Silva and da Cruz e Silva 2003), the A β PP⁷⁵¹ isoform is most abundant in peripheral tissues (Tanaka *et al.* 1989). In s-IBM muscle fibers, A β PP⁷⁵¹ is the isoform overproduced and accumulated as aggregates (Askanas *et al.* 1993; Sarkozi *et al.* 1993; Guerin *et al.* 2008). Phosphorylation of A β PP is considered a regulatory mechanism of A β PP metabolism (da Cruz e Silva and da Cruz e Silva 2003). Phosphorylation on Threonine 668 of the neuronal isoform A β PP⁶⁹⁵ (equivalent to threonine at position 724 of A β PP⁷⁵¹) was reported a) to be associated with increased A β production (Lee *et al.* 2003; Ando *et al.* 2001), and b) to mediate pathological interaction between A β and tau (Shin *et al.* 2007). There were reduced levels of A β in neuronal cells treated with T688 kinase inhibitors or overexpressing mutant p-A β PP^{T688} (Lee *et al.*, 2003).

Active p-GSK3 β ^{Y216} has been shown increased in the frontal cortex of AD patients (Leroy *et al.* 2007), and proposed to be a component of Alzheimer disease pathogenesis (Jope 2003; Jope and Johnson 2004). Active p-GSK3 β ^{Y216} was shown to phosphorylate both A β PP on Threonine 688 (Aplin *et al.* 1996) and tau protein (Anderton *et al.* 2001).

GSK3 β is inactivated by being phosphorylated on Serine 9 (Bhat *et al.* 2000; Shaw *et al.* 1997). In AD transgenic mice, overexpression of a dominant negative GSK3 β , as well as treatment with lithium or other pharmacological inhibitors, was reported to result in GSK3 β inhibition and a subsequent reduction of A β , which was associated with improved cognitive performance (Rockenstein *et al.* 2007; Ryder *et al.* 2003; Su *et al.* 2004). GSK3 β activity was also found increased in muscle fibers of the transgenic-mouse IBM model (Kitazawa *et al.* 2006); in this model, a decrease of GSK3 β activity by lithium correlated with decreased tau phosphorylation (Kitazawa *et al.* 2008). In the present study, we demonstrated for the first time that a) in biopsied s-IBM muscle fibers and in our A β PP+ cultured human muscle-fibers (A β PP+ human muscle culture IBM model), A β PP is phosphorylated on Threonine⁷²⁴, and b) in s-IBM patients, active GSK3 β is significantly increased as compared to the normal aged-matched-control muscle biopsies. In addition, we demonstrated in our IBM culture model that proteasome inhibition significantly enhanced the increase of active GSK3 β , which corresponded to the increase of phosphorylated A β PP. Accordingly, we postulate that in s-IBM, phosphorylation of A β PP is influenced by proteasome inhibition, possibly via activation of GSK3 β . The increase of total A β PP after proteasome inhibition also suggests that in this model the ubiquitin-proteasome system is involved in the degradation of A β PP.

Our results showed that lithium increased activities of all three proteasome enzymes studied, to various degrees. Lithium is known to have various effects on cells (Phiel and Klein 2001), but to our knowledge it has not been reported to improve proteasome function. However, since A β PP was previously shown to inhibit proteasome function in this culture model (Fratta *et al.* 2005), it is presently not known whether the increase of proteasome function by lithium that we observed in the current study represents its direct influence on the ubiquitin-proteasome system, or it is related to the overall reduction of the A β PP level.

In conclusion, we have demonstrated, apparently for the first time, that in s-IBM muscle fibers GSK3 β activity is increased and A β PP is phosphorylated. Our experimental data also

strongly suggest that A β PP phosphorylation is increased by GSK3 β activation, which is increased by proteasome inhibition.

And, we have shown that in our culture IBM model, lithium treatment significantly decreased the levels of both total and phosphorylated A β PP as well as A β oligomers, accompanied by increased proteasome function and decreased GSK3 β activation. This is in agreement with the transgenic-mouse IBM model in which lithium treatment decreased both GSK3 β activation and tau phosphorylation (Kitazawa *et al.* 2008). Accordingly, treatment with lithium, or other GSK3 β inhibitors, might be beneficial for s-IBM patients.

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Abbreviations used

Aβ	amyloid- β
AβPP	amyloid- β precursor protein
AD	Alzheimer disease
ALS	amyotrophic lateral sclerosis
CHMFs	cultured human muscle fibers
CTL	chymotrypsin-like
GSK-3β	glycogen synthase kinase 3 β [E.C. 2.7.11.36]
Epox	epoxomicin
ER	endoplasmic reticulum
PGPH	peptidyl -glutamyl peptide hydrolyzing
S (Ser)	Serine
s-IBM	sporadic inclusion-body myositis
T (Thr)	Threonine
TL	trypsin-like
Tm	tunicamycin
Tg	thapsigargin
Y (Tyr)	Tyrosine

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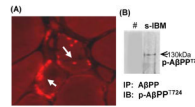


Fig. 1.

Phosphorylated AβPP T724 (p-AβPP^{T724}) in sporadic-inclusion body myositis (s-IBM) muscle fibers. (A) Immunofluorescence using the specific phospho-AβPP antibody illustrates pAβPP^{T724}-immunoreactive aggregates in two muscle fibers (arrows). (B). Immunoprecipitation (IP) of p-AβPP^{T724} in s-IBM muscle biopsy. IP was performed using 6E10 antibody, which recognizes both AβPP and Aβ; immunoblots (IB) were subsequently probed with the specific anti p-AβPP^{T724} rabbit polyclonal antibody. There is a distinct band corresponding to AβPP^{T724}.

In # the primary antibody was omitted from the immunoprecipitation reaction to determine the specificity of the reaction.

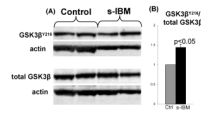


Fig. 2.

GSK3 β is activated in s-IBM biopsied muscle. (A) Representative immunoblots of normal control and s-IBM muscle biopsies. (B) The ratio of GSK3 β ^{Y216} to total GSK3 β , obtained by densitometric analysis of protein bands, shows that in s-IBM, as compared to controls, GSK3 β ^{Y216} is significantly increased ($p < 0.05$, 2 tail Student's t-test). \pm SEM.

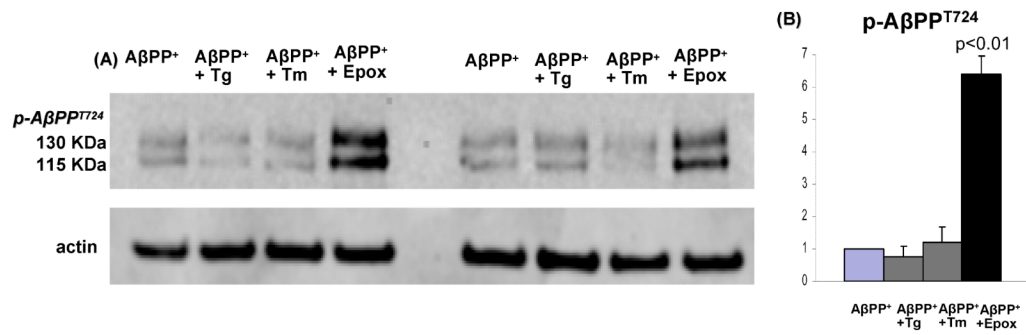


Fig. 3. p-AβPP^{T724} in i) proteasome-inhibited, and ii) ER-stress-induced human-muscle-culture AβPP⁺ IBM model. (A) Representative immunoblots of densitometric analysis in (B). (B) Densitometric analysis based on 12 independent experiments of the pAβPP^{T724} protein bands relative to the actin bands, expressed in arbitrary units, shows that after treatment with Epoxomicin (Epox) pAβPP^{T724} is significantly increased (6.5 fold, $p < 0.01$, 2 tail Student's t-test), while the ER stress inducers (Thapsigargin [Tg] and Tunicamycin [Tm]) did not exert this effect. \pm SEM.

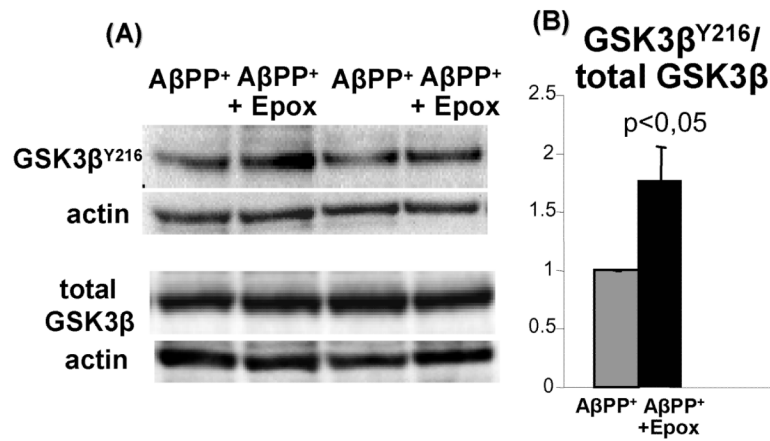


Fig.4. GSK3β is activated in the epoxomicin-inhibited proteasome in the human-muscle-culture IBM model. (A) Representative immunoblots of densitometric analysis in (B). (B) The ratio of GSK3β^{Y216} to total GSK3β obtained by densitometric analysis of protein bands in 6 independent experiments shows that in proteasome-inhibited cultures, as compared to controls, GSK3β^{Y216} is significantly increased (1.7 fold, $p < 0.05$, 2 tail Student's t-test) \pm SEM.

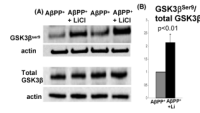


Fig.5.

Inactive form of GSK3 β is increased in LiCl-treated culture-IBM-model. (A) Representative immunoblot of GSK3 β ^{Ser9} in LiCl-treated and sister-control untreated culture A β PP⁺ IBM model. (B) Densitometric analysis of GSK3 β ^{Ser9} bands relative to the actin bands, expressed in arbitrary units, shows that after lithium treatment GSK3 β ^{Ser9} is significantly increased (2.2, $p < 0.01$, 2 tail, Student's t-test).

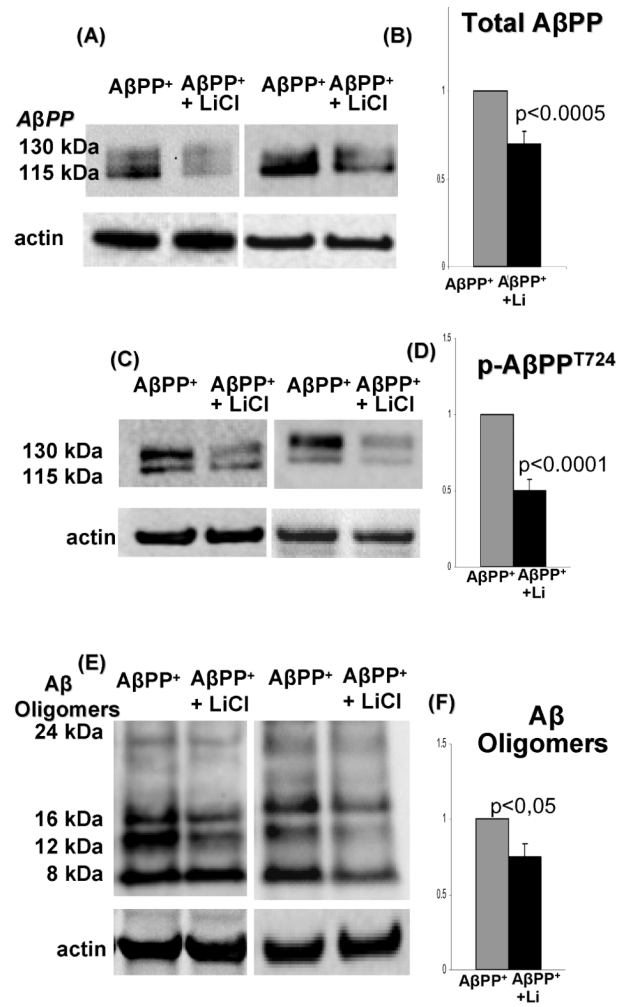


Fig. 6. Lithium decreases the amount of total AβPP, p-AβPP^{T724}, and Aβ-oligomers. (A) Representative immunoblots of total AβPP in LiCl-treated and control untreated culture AβPP⁺-IBM-model. (B) Densitometric analysis of the total AβPP bands (130 and 115 kDa) relative to the actin bands, expressed in arbitrary units, shows that after lithium treatment AβPP is decreased 30% (p < 0.0005, 2 tail); (C) Representative immunoblots of p-AβPP^{T724} in LiCl-treated and untreated control culture AβPP⁺-IBM-model. (D) Densitometric analysis of p-AβPP^{T724} bands relative to the actin bands, expressed in arbitrary units, shows that after lithium treatment p-AβPP^{T724} is decreased 50% (p < 0.0001, 2 tail). (E) Representative immunoblots of Aβ oligomers (8 kDa, 12 kDa, 16 kDa, 25 kDa bands) in LiCl-treated and control untreated AβPP⁺-IBM-culture-model. (F) Densitometric analysis of Aβ oligomers bands relative to the actin bands, expressed in arbitrary units, shows that after lithium treatment all Aβ bands calculated together were decreased 25% (p < 0.05, 2 tail Student's t-test).

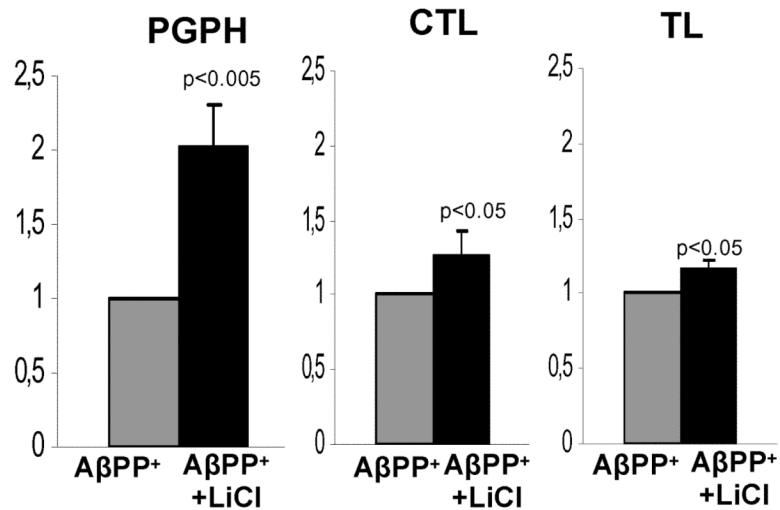


Fig.7.

Three main proteasome enzyme activities -- peptidyl-glutamyl-peptide hydrolytic (PGPH), chymotrypsin-like (CTL) and trypsin-like (TL) -- in LiCl treated and sister-control untreated culture AβPP+ IBM model. Proteasome activities are expressed per β2 20S proteasome subunit in each culture. After lithium treatment, PGPH activity was increased 2 fold ($p<0.005$, 2 tail Student's t-test), CTL activity was increased 1.3 fold ($p<0.05$, 1 tail Student's t-test), and TL activity was increased 1.2 fold ($p<0.05$, 1 tail Student's t-test).