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α-synuclein aggregates with β-amyloid or tau in human red blood cells: correlation with antioxidant capability and physical exercise in human healthy subjects --Manuscript Draft--

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Abstract:	Neurodegenerative disorders (NDs) are characterized by abnormal accumulation/misfolding of specific proteins, primarily α -synuclein (α -syn), β -amyloid 1-42 (A β) and tau, in both brain and peripheral tissue. In addition to homo-oligomers, the role of α -syn interactions with Ab or tau has gradually emerged. The altered protein accumulation has been related to both oxidative stress and physical activity; nevertheless, no correlation among the presence of peripheral a-syn hetero-aggregates, antioxidant capacity and physical exercise has been discovered as of yet. Herein, the content of α -syn, A β , tau and of their heterocomplexes was determined in red blood cells (RBCs) of healthy subjects (sedentary and athletes). Such parameters were related to the extent of the antioxidant capability (AOC), a key marker of oxidative stress in aging-related pathologies, and to physical exercise, which is known to play at important preventive role in NDs and to modulate oxidative stress. Tau content and plasma AOC toward hydroxyl radicals were both reduced in older or sedentary subjects; in contrast, α -syn and A β accumulated in elderly subjects and showed an inverse correlation with both hydroxyl AOC. Furthermore, α -syn/A β aggregates were significantly reduced in athletes and inversely correlated with physica activity level, independent of age. The positive correlation between antioxidant capability/physical activity and reduced protein accumulation was confirmed by these data, and suggest that peripheral α -syn heterocomplexes may represent new indicators of ND-related protein misfolding.



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Executive Editor

Molecular Neurobiology

Dear Editor,

Herewith enclosed please find the revised version of our manuscript entitled " α -synuclein aggregates with β -amyloid or tau in human red blood cells: correlation with antioxidant capability and physical exercise in human subjects" by Daniele S. et al.

We are grateful for the opportunity to revise our manuscript and to improve its quality by following the referee's suggestions. We do hope the new version of the manuscript will meet all the requirements for publication in Molecular Neurobiology

Looking forward to hearing from you soon, we thank you and send you our best personal regards. Sincerely yours,

Maria Letizia Trincavelli and Ferdinando Franzoni



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Manuscript #MOLN-D-17-00041.

Title:

Major Points

1. Some of the parameter is difficult to be understood. Firstly, why the tau protein in young SED is higher than all other group including young ATHL and old SED which seemed to be in opposite population. This data suggested that ATHL has no effect with physical exercise.

As reported in Table 2, the RBC tau concentrations were $10,51\pm8,34$ ng/mg protein and $6,67\pm3,28$ ng/mg protein in young and old SED, respectively, suggesting that tau does not accumulate with age, at least in the analysed population. Consistent with these data, RBC levels of total tau significantly decreased with age in the total population (Fig. 4d, young versus older, P=0.0434). Further studies will be needed to clarify the significance of tau decrease with age, also considering that literature data are not available for this parameter in RBCs.

Moreover, tau levels in RBCs were demonstrated to be almost the same levels in ATHL $(6,32\pm3,67$ ng/mg protein and $5,59\pm4,12$ ng/mg protein in young and old ATHL, respectively). Based on these findings, it could be speculated that physical exercise can exert beneficial effects lowering tau concentration in particular in the young subgroup (see Figure 4d).

To clarify the data obtained on tau levels in RBCs, results and discussion concerning tau levels were modified in the text (see Results section, page 14, and Discussion section, page 22).

2. Correlation data showed in Fig 5 and Fig 6 lack some of the parameter/ Author should show all the correlation form each populations.

As required by the referee, all the correlations were added in the revised version of the manuscript (see the new Figures 5 and 6, and the new Supplementary Figures 1, 2 and 3). The data were omitted in the first version of the manuscript due to the lack of significant change in the parameters.

3. Authors detected misfolding proteins from RBC. Clarify whether excise could alter the RBC turnover that can effect on the expression of those proteins. Authors should measure the turnover of red blood cells and compared that between ATHL and SED groups.

We thank the referee for his interesting comment. Intensive training could affect RBC turnover (Sports Med. 1995;19(1):9-31) and reduce RBC mass by intravascular hemolysis of senescent red blood cells. (Front Physiol. 2013; 4: 332). Actually, conflicting data have been emerging (Adv Clin J Lab Hematol. Chem. 2013: 59:125-53: Int 2011 Dec: 33(6):638-4; Acta Haematol. 2012;127(3):156-64), indicating that RBC production in athletes does not significantly differ from sedentary controls. Besides the actual effects of physical training, measuring RBC turnover in our cohort would require an ex novo recruitment of subjects and blood collection. For this reason, the influence of exercise on RBC ageing will be take into consideration in a future study. In the meantime, such criticism was discussed in the revised version of the manuscript (see Discussion section, page 23).

4. The number of samples is too small to represent WB and immunoprecipitation figures. Authors should add samples more to make quantitative figures.

Western blot analyses were conducted to qualitatively demonstrate that the analysed proteins (i.e. α -syn, A β , α -syn/A β , tau and α -syn/tau) were expressed in RBCs (see Figure 1). Moreover, such assay was utilized to show the physical interaction *in vitro* of α -syn with A β or tau (see Figure 2). In contrast, no comparison between groups was conducted using western blot data. In order to avoid misunderstanding, this issue was clarified in the revised version of the manuscript (see Methods section, page 6).

5. Have authors ever tried to detect the level of those misfolding proteins to see whether they have any correlation of those expression in red blood cells.

We thank the referee for his suggestion. In this paper, the levels of the oligomeric form of α -syn were measured in RBCs. Unfortunately, no significant differences between SED and ATHL were found (see Figure 4B). As concern the other proteins, new specific antibodies and techniques have been emerging to detect oligomeric forms of tau (*FASEB J 2012; 26(5): 1946–1959; Front Neurol. 2014; 5: 251*) and A β (*Cell Rep. 2014;7(1):261-8;J Neurosci. 2014 Feb 19;34(8):2884-97*). We agree with the referee that measuring the misfolded forms of ND-related proteins will be an interesting point to assess the contribution of both oxidative stress and physical exercise, thus these points were added in the text as possible future goals (See Discussion section, page 23).

6. In figure 1b, there are bands which do not indicate A β (middle panel: more than 25kDa) or tau (lower panel: less than 50kDa). What do these bands mean?

Bands with a higher molecular weight than 25kDa (Fig. 1b, middle panel) could indicate elevated oligomeric form of A β (*JBC*, 2015; 290: 17415–17438; *PLoS One*.2014;9:e114041). In contrast, bands lower than 50 kDa (Fig. 1b, lower panel) have been related to truncated or cleaved forms of tau (*Neuroscience Letters 2006; 399:106–110*). Such findings were reported in the revised version of the manuscript (Results section, page 11).

7. In the result, the authors showed that hydroxyl TOSC values were inversely correlated with RBC total α -syn in the ATHL subgroup. Is there any correlation between hydroxyl TOSC and total α -syn concentration in SED group?

At P16 line3, authors concluded that α -syn accumulation in RBCs is strongly related to hydroxyl AOC. There is no significant correlation between α -syn concentration and hydroxyl AOC in young subgroup. Authors should try to discuss the discrepancy.

Any significant correlation between hydroxyl TOSC and total α -syn concentration was found in SED group (P=0.223, R²=0.0097, new Suppl. Fig. 1a), although a trend toward an inverse correlation was evidenced. These findings were added in the revised version of the manuscript (Results section, page 16).

Since the correlation between hydroxyl TOSC and total α -syn concentration was not found in the young subgroup, as well as in the SED cohort, the sentence reported at page 16 was modified. In particular, based on our results, a higher contribute of hydroxyl AOC on α -syn accumulation can be speculated with advanced age or physical exercise (see Results section, page 16).

8. In figure 7, the authors should add the scatter plot of SED group. The scatter plot of hydroxyl TOSC vs α -syn/A β levels of the SED group was added in the manuscript (see new Figure 7e).

In figure 8, the authors should add the scatter plot of all groups (Older, Young, ATHL, SED group). The scatter plots of hydroxyl TOSC vs α -syn/tau levels of the missing groups were added in the manuscript (see new Figure 8d and e, and new Suppl. Fig. 4).

3

In figure 9, the authors should add the plot of all groups (Older, Young group) about all proteins. The scatter plot of all the groups were added in the new version of the manuscript (see new Supplementary Figures 5-7).

Minor Points

- 1. Authors mistake numerical display in Table 1 and 2. For example, 37,50. Numerical display in Table 1 and 2 were corrected. Thank you.
- "e4 allele" in figure legend (P25-Line27) should be "ε4 allele." The figure legend was corrected. Thank you.
- 3. "e4 allele" in figure legend(P26-Line2) should be "ε4 allele." The figure legend was corrected. Thank you.
- 4. Authors mentioned they used A β antibody in section "Co-immunoprecipitationwestern blotting". However, they showed A β 1-42 in Fig1. Which is correct? A β antibody in section "Co-immunoprecipitation-western blotting" was corrected in A β 1-42.

 α -synuclein aggregates with β -amyloid or tau in human red blood cells: correlation with antioxidant capability and physical exercise in human healthy subjects

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Abstract

Neurodegenerative disorders (NDs) are characterized by abnormal accumulation/misfolding of specific proteins, primarily α -synuclein (α -syn), β -amyloid₁₋₄₂ (A β) and tau, in both brain and peripheral tissue. In addition to homo-oligomers, the role of α -syn interactions with A β or tau has gradually emerged. The altered protein accumulation has been related to both oxidative stress and physical activity; nevertheless, no correlation among the presence of peripheral α -syn hetero-aggregates, antioxidant capacity and physical exercise has been discovered as of yet.

Herein, the content of α -syn, A β , tau and of their heterocomplexes was determined in red blood cells (RBCs) of healthy subjects (sedentary and athletes). Such parameters were related to the extent of the antioxidant capability (AOC), a key marker of oxidative stress in aging-related pathologies, and to physical exercise, which is known to play an important preventive role in NDs and to modulate oxidative stress.

Tau content and plasma AOC toward hydroxyl radicals were both reduced in older or sedentary subjects; in contrast, α -syn and A β accumulated in elderly subjects and showed an inverse correlation with both hydroxyl AOC and the level of physical activity. For the first time, α -syn heterocomplexes with A β or tau were quantified and demonstrated to be inversely related to hydroxyl AOC. Furthermore, α -syn/A β aggregates were significantly reduced in athletes and inversely correlated with physical activity level, independent of age.

The positive correlation between antioxidant capability/physical activity and reduced protein accumulation was confirmed by these data, and suggest that peripheral α -syn heterocomplexes may represent new indicators of ND-related protein misfolding.

Key words: protein misfolding; neurodegenerative diseases; α -synuclein; β -amyloid; tau; α -synuclein heterocomplexes; antioxidant capability; physical exercise.

Introduction

Neurodegenerative disorders (NDs), such as Alzheimer's disease (AD), Parkinson's disease (PD) and amyotrophic lateral sclerosis, are characterized by the pathological evidence of progressive neuronal loss in specific vulnerable areas [1-3] and reactive gliosis [4, 5], in which self-aggregating misfolded proteins form high-ordered insoluble fibrils in neurons and/or glial cells [4-6]. For example, abnormal accumulation of α -synuclein (α -syn) has been established to form Lewy bodies and neurites in PD and dementia with Lewy bodies (DLB), as well as glial cytoplasmic inclusions in multiple system atrophy [7]. In contrast, the neuropathological hallmarks of AD are represented by senile plaques containing extracellular deposits of β -amyloid₁₋₄₂ (A β) and by intra-neuronal neurofibrillary tangles composed of hyperphosphorylated tau protein [8, 9]. However, postmortem evaluation often discloses a mixed pattern of proteinopathies, commonly accompanied by signs of chronic cerebrovascular disease pathology. In this regard, the potential contribution of α -syn to AD pathogenesis is emerging as well [10, 11], with 30–40 % of AD cases presenting Lewy bodies and Lewy neurites [12].

Data from the literature suggest that $A\beta$, tau and α -syn might promote the accumulation or aggregation of one another [13]. Furthermore, in addition to homo-oligomers, the role of heterocomplexes has been emerging; $A\beta$ and α -syn have been shown to co-immunoprecipitate and form complexes in patient brains and transgenic models, providing clear evidence for their direct interaction [14, 15]. Additionally, in different cellular systems, α -syn has been demonstrated to bind to tau directly and promote the polymerization of the latter [13, 16-18], even in the axonal compartment [19]. Of note, the alterations in protein misfolding related to NDs are not restricted within the brain, but appear in peripheral tissue as well. For this reason, great efforts have been devoted to exploiting substantial biological changes and putative biomarkers in tissue other than the brain or cerebrospinal fluid (CSF) [20-24]. In this respect, the use of blood has gradually emerged due to its availability, low cost and time effectiveness [25].

ND-protein oligomerization has been linked not only to genetic factors but also to environmental factors [26, 27], primarily including oxidative stress [26, 28], which arises from an imbalance between an excessive generation of reactive oxygen species (ROS) and the biological system's capability to eliminate the reactive intermediates [29]. Oxidative stress leads to a progressive decline in cell physiology [30] by damaging cellular macromolecular components such as DNA, lipids and proteins [31, 32]. As oxidative stress strongly contributes to abnormal protein misfolding and the propagation of oligomers, antioxidant therapies are emerging as a preventive therapeutic option for age-related NDs [33, 34].

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Epidemiological studies have extensively demonstrated that regular exercise is an important preventive and therapeutic tool in AD, PD and cerebrovascular diseases [35]. The adaptive response to regular exercise involves the up-regulation of the enzymatic antioxidant system and modulation of oxidative damage, which culminates in a recovered redox state of brain cells [36, 37]. In addition to a reduction in oxidative damage, the effects of exercise seem to be quite intricate, including an increase in neurogenesis and capillarization and an enhanced proteolytic degradation of toxic oligomers by proteasomes and neprilysin [38-40]. In a recent paper, voluntary running has been shown to counteract amyloid deposition, tau phosphorylation, inflammatory reaction, and memory loss in a mouse model of AD [41]. Moreover, lower $A\beta$ plasma concentrations and brain depositions have been observed in humans performing higher levels of physical activity [42], consistent with the hypothesis that exercise may be involved in the modulation of pathogenic changes associated with NDs [43].

All together, these data highlight the link between neurodegeneration, oxidative stress and physical exercise. However, the correlation among antioxidant capability, physical exercise and the accumulation of α -syn heterocomplexes with tau and A β at a peripheral level has never been investigated. The aim of the current study was to examine how the levels of protein aggregates, associated with ND development, could be influenced by the intrinsic plasma antioxidant capacity and by physical activity. More specifically, plasma antioxidant capability (AOC), the primary marker of oxidative stress in aging-related pathologies [44], was measured in a cohort of young and older athletes and sedentary subjects, and related to the grade of physical exercise. Moreover, the content of total and oligomeric α -syn, tau and A β was revealed in red blood cells (RBCs) of the same subjects. Indeed, these cells represent a good model to study the aging-related biochemical alterations, including protein misfolding [42, 45, 46]. Furthermore, α -syn heterocomplexes with A β or tau were detected and quantitatively measured to unveil their presence and putative role as novel indicator of oxidative stress-related neurodegeneration in peripheral fluids.

Materials and Methods

Ascorbic acid, 2,20-azo-bisamidinopropane (ABAP), Iron (III) chloride hexa-hydrate, KMBA (acheto-g-(methylthiol)butyric acid) and recombinant human α -syn, tau and A β were purchased from Sigma Chemical Co. (St. Louis,MO, USA). Antibodies to α -syn, tau and A β were from Santa Cruz Biotechnology.

Study population and setting of the study

Forty-eight sex-matched endurance athletes (ATHL, mean age 44.6±13.4 years, range 22-74 years) recruited from the Sport Medicine Unit of the Department of Clinical and Experimental Medicine of the University of Pisa, and 58 healthy age-sex-matched sedentary volunteers (SED, mean age 46.7±14.5 years, range 20-75 years) were studied (Table 1). Athletes performed endurance exercise more than three times/week and were active in national road-running races.

All subjects were free of cardiovascular disease or other major medical disorders, as assessed by clinical history, physical examination, basal and stress electrocardiography, blood chemistry, hematology and urine analysis. Major criteria for inclusion of subjects in the trial were as followed: total plasma cholesterol ranging from 3.1 to 5.8 mmol/L, HDL cholesterol from 0.67 to 1.9 mmol/L, plasma triglycerides from 0.34 to 1.7 mmol/L, body mass index lower than 30 kg/m², diastolic arterial blood pressure lower than 90 mmHg and systolic arterial blood pressure lower than 140 mmHg. Subjects were excluded if they had smoking habits or received any drug treatment within the previous two months.

Aerobic fitness was evaluated with a maximal graded cycle ergometry test performed by a cardiologist blinded to the other data. Participants started at 25 Watts. Increments of 25 watts per 2 min were made until exhaustion or until reaching one of the American College of Sports Medicine established criteria for maximal oxygen uptake [47]. Heart rate was continuously measured by ECG. Blood pressure and the rate of perceived exertion (RPE) were assessed at the end of each step. Recovery was monitored until heart rate was <100 bpm. The maximum achieved resistance (Watts) was retained for all calculations. The 15-point Borg RPE scale [48, 49] was used to evaluate the level of intensity for each participant. The scale ranges from 6 to 20, with 6 corresponding to no exertion at all, 7.5 to extremely light, 9 to very light, 11 to light, 13 to somewhat hard, 15 to hard, 17 to very hard, 19 to extremely hard, and 20 to maximal exertion.

Each group was divided into a younger (<50 years) and an older (>50 years) subgroup (Table 1). The time period between the last exercise bout and blood sampling was at least 48 hours.

This study was approved by the Ethics Committee of the Great North West Area of Tuscany (271/2014 to F.F.) and it was carried out in accordance with the Declaration of Helsinki. All

subjects gave informed consent to participate in the study. Fully informed consent was obtained from each subject entering the study.

AOC was measured in plasma samples from the aforementioned cohort. RBCs were collected to quantify the content of total and oligomeric α -syn, tau, A β and α -syn heterocomplexes with A β or tau.

Plasma and RBC collection.

Whole blood was collected from healthy volunteers (N=106, Table 1) into a tube containing EDTA as an anticoagulant. RBCs were separated from plasma by a centrifugation at $200 \times g$ at 4 °C for 10 minutes. The plasma was then stored at -80°C in different aliquots. The RBC pellet was centrifuged at 1000 x g for 10 min and washed three times with PBS. RBC pellet was frozen at -20 °C until use.

Co-immunoprecipitation–western blotting

In order to verify α -syn, tau and A β expression in human blood, RBCs (50 mg) were lysed with RIPA buffer [50] and then resolved by SDS-PAGE (8.5%). Samples were probed overnight at 4°C with primary antibodies to α -syn (α/β -synuclein N-19, SC-7012, Santa Cruz Biotechnology), tau (H-150 SC-5587, Santa Cruz Biotechnology) or A β_{1-42} (β -amyloid H-43 SC-9129, Santa Cruz Biotechnology). The primary antibodies were detected using peroxidase- conjugated secondary antibodies and a chemioluminescent substrate (ECL, Perkin Elmer).

To confirm the presence of α -syn heterocomplexes with tau or A β , a co-immunoprecipitation assay was employed [50]. Briefly, 1 mg of lysates obtained from RBCs was resuspended in RIPA buffer and was probed overnight under constant rotation with an anti- α -syn antibody (5 µg/sample), and then immunoprecipitated with protein A-Sepharose. After extensive washing, the immunocomplexes were resuspended in Laemmli solution, resolved by SDS-PAGE and probed overnight with primary antibodies to α -syn (input), tau or A β_{1-42} as described above.

Of note, western blot analyses were used as qualitative data on ND-related protein expression in RBCs.

Immunoassay methods for total a-synuclein

Total α -synuclein was detected in RBCs following literature's protocols [51]. Briefly, wells were pre-coated overnight at 4°C with a full length polyclonal antibody to α -syn (sc-10717, Santa Cruz Biotechnology), and non-specific sites were blocked using bovine serum albumine (BSA) for 1 h at 37°C. RBCs (0,150 mg/100 µl) were captured on wells for 2h at 25°C. Purified recombinant protein standards of α -syn were assayed in parallel with human samples to generate a standard curve. After

extensive washing, samples were probed with a mouse monoclonal antibody to α -syn (Santa Cruz, sc-12767), and subsequently with an anti-mouse-HRP antibody. The wells were then washed 4 times with PBS-T (phosphate buffered saline containing 0.01% Tween 20), before adding the enzyme substrate TMB (3,3',5,5'-tetramethylbenzidine, Thermo Scientific) and leaving the colour to develop for 30 min at room temperature. Absorbance values at 450 nm.

Preparation of aged solutions of α -syn and of the α -syn biotinylated antibody

Recombinant α -syn were incubated in parafilm-sealed tubes at 37°C for 4 days in an Eppendorf Thermomixer with continuous mixing (1000 rpm), as reported previously [52].

To prepare the α -syn biotinylated antibody, Sulfo-NHS-LC-Biotin (Pierce, Rockford, IL, USA) (200 mg) was reacted with the 211 mouse monoclonal antibody (mAb) (Santa Cruz Biotechnology, Santa Cruz, CA, USA) [53]. The mixture was desalted on Bio-Spin-6 columns (BIO-RAD, UK) to remove excess uncoupled biotin.

Detection of oligomeric a-syn

Oligomeric α -syn levels in RBCs were measured using an immunoenzymatic assay, as previously described [51, 52]. The plate was pre-coated overnight at room temperature with the mouse monoclonal α -syn 211 antibody (Santa Cruz, sc-12767). After extensive washing with PBS-T, non-specific sites were blocked with 1% BSA. RBCs (0,04 mg/100 µl) were added to each well and incubated at 25°C for 2 h. α -syn oligomers were detected using the α -syn biotinylated antibody, which recognizes amino acid residues 121-125 of human α -syn. Whereas for antigen detection, streptavidin-horseradish peroxidase conjugate antibody (1:1000, GE Healthcare) was used. The wells were washed three times with PBS-T, before the addition of 100 µl/well of TMB, as reported above.

Detection of total Aß

A β levels in blood samples were measured using an immuno-enzymatic assay, as described previously [53]. The plate was pre-coated overnight at 4°C with a specific antibody to A β (Santa Cruz, sc-9129). After extensive washing with PBS-T, non-specific sites were blocked with 1% BSA. RBCs (0,2 mg/100 µl) were added to each well and incubated at 25°C for one hour. After extensive washing with PBS-T, samples were detected using the polyclonal antibody to A β (sc-5399, Santa Cruz Biotechnology). The standard curve was constructed using recombinant human A β solutions at eight different concentrations.

Detection of total tau

Tau levels in blood samples were measured using an immuno-enzymatic assay, as described previously [53]. The plate was pre-coated overnight at 4°C with a specific antibody to tau (Santa Cruz, sc-32274). After extensive washing with PBS-T, non-specific sites were blocked with 1% BSA. RBCs (0,5 mg/100 μ l) were added to each well and incubated at 25°C for one hour. After extensive washing with PBS-T, samples were detected using the polyclonal antibody to tau (sc-5587, Santa Cruz Biotechnology). The standard curve was constructed using recombinant human tau solutions at eight different concentrations.

Immunoassay detection of α-syn/Aβ heterocomplexes

For the quantification of α -syn/A β interactions, a 'home-made' method was developed employing a "sandwich" immunoenzymatic assay [54, 55], as follows. Standard α -syn/A β were prepared by incubating 1 mg of each protein, diluited in 2 mM sodium dodecyl sulfate (SDS) in parafilm-sealed tubes at 37°C for 36 h in an "Eppendorf Thermomixer" with continuous mixing (500 rpm) [14]. Eight different dilutions of α -syn/A β were prepared; following capturing on wells pre-coated with a specific antibody to A β (see below), the samples were collected, and α -syn levels (i.e., the quotes of recombinant α -syn not bound to A β) were quantified by a validated immunoenzymatic assay described above.

The microplate was pre-coated overnight at room temperature with β -amyloid H-43 antibody (1:100, sc-9129, Santa Cruz Biotechnology) in poli-L-ornithine/NaHCO₃, pH 9.6. After two washes with PBS-T, RBCs (40 mg/sample in 2 mM SDS) were added to each well and incubated at 25°C for 2h. The wells were washed, and non-specific sites were blocked with 1% BSA for 30 min at 37°C. To detect α -syn bound to A β , samples were probed for 2 h 37°C with a specific antibody to α -syn (sc-12767, Santa Cruz Biotechnology), and subsequently with the appropriate HRP-conjugated antibody. After 1.5 h, the wells were washed twice with PBS-T, before the addition of 100 µl/well of TMB. Absorbance was measured at 450 nm. Relative concentration of α -syn/A β complexes were calculated according to the standard curve obtained in each microplate. The assays of blood plasma were all carried out in duplicate. Blood samples from athletes and sedentary subjects were analysed together in batch runs. For some subjects, multiple assays were performed on diluted RBCs from a single subject to confirm that low or high concentrations were in the linear range of the assay. All measurements were repeated twice and the average value was determined.

Immunoassay detection of α-syn/tau heterocomplexes

For the quantification of α -syn/tau interactions, a similar 'home-made' method was developed (see the precedent paragraph). Standard α -syn/tau were prepared by incubating 1 mg of each protein, diluted in 2 mM SDS in parafilm-sealed tubes at 37°C for 1 h in an Eppendorf Thermomixer with continuous mixing (500 rpm). Following capturing on wells pre-coated with a specific antibody to α -syn (see below), the samples were collected, and tau levels (i.e., the quotes of recombinant α -syn not bound to A β) were quantified by a validated immunoenzymatic assay described above.

The microplate was pre-coated overnight at room temperature with anti- α -syn antibody (1:100, sc-7012, Santa Cruz Biotechnology) in poli-L-ornithine/NaHCO₃, pH 9.6. After two washes with PBS-T, RBCs (80 mg/sample in 2 mM SDS) were added to each well and incubated at 25°C for 2h. The wells were washed, and non-specific sites were blocked with 1% BSA for 30 min at 37°C. To detect α -syn bound to tau, samples were probed for 2 h 37°C with a specific antibody to tau (sc-5587, Santa Cruz Biotechnology), and subsequently with the appropriate HRP-conjugated antibody. After 1.5 h, the wells were washed twice with PBS-T, before the addition of 100 µl/well of TMB. Absorbance was measured at 450 nm. Relative concentration of α -syn/tau complexes were calculated according to the standard curve obtained in each microplate.

Total Oxyradical Scavenging Capacity (TOSC) Assay

The plasma antioxidant capability (AOC) was assessed by the TOSC assay, a gas chromatographic assay for determining oxyradical scavenging capacity of biological fluids [56, 57]. Peroxyl radicals were generated by thermal homolysis of 20 mM ABAP at 35°C in 100 mM potassium phosphate buffer, pH 7.4. Hydroxyl radicals were generated at 35°C by the iron plus ascorbate-driven Fenton reaction (1.8 mM Fe3+, 3.6 mM EDTA, and 180 mM ascorbic acid in 100 mM potassium phosphate buffer, pH 7.4). Reactions with 0.2 mM KMBA were carried out in 10 ml vials sealed with gas-tight Mininert1 valves (Supelco, Bellefonte, PA) in a final volume of 1 ml.

Ethylene production was measured by gas-chromatographic analysis of 200 µl aliquots taken from the headspace of vials at timed intervals during the course of the reaction. Analyses were performed with a Hewlett-Packard gas chromatograph (HP 7820A Series, Andoven, MA) equipped with a Supelco DB-1 (30 x 0.32 x 0.25 mm) capillary column and a flame ionization detector (FID). The oven, injection and FID temperatures were respectively 35,160 and 220°C. Hydrogen was the carrier gas (flow rate of 1 ml/min) and a split ratio of 20:1 was used. Total ethylene formation was quantified from the area under the kinetic curves that best define the experimental points obtained for control reactions and after addition of plasma during the reaction [56-58]. TOSC values were quantified from the equation TOSC = 100 - (SA/CA x 100), where SA and CA are respectively the area under the curve (AUC) for sample and control reaction. A TOSC value of 0 corresponds to a

sample with no scavenging capacity. A TOSC value of 100 is attributed to a compound that entirely suppresses the ethylene formation whereas a pro-oxidant compound shows a negative TOSC value [59]. Consequently, antioxidants and pro-oxidants molecules can be distinguished by the obtained results. The linearity of dose–response curve between plasma (ml) and the antioxidant response (TOSC value) was tested and good correlation coefficients (generally greater than 0.9) were obtained at the different doses used to test the validity of our experiments. Each experiment was performed in duplicate to account for the intrinsic variability of the method. The results obtained with plasma were expressed in TOSC units. In our hands, the coefficient of variation (CV) of the method ranged between 2% and 5% [57, 60].

Genotyping

Apolipoprotein E (Apo E) genotypes were identified by restriction fragment length polymorphism (RFLP), using genomic DNA extracted from blood of heatlhy subjects (N=87, Table 1). PCR was performed using 1.5 pmol of each primer (forward: 5 ' -TCG-GCCGCA-GGG-CGC-TGA-TGG-3 ' and reverse: 5 ' -CTCGCG-GGC-CCC-GGC-CTG-GTA-3 '), 250 µmol/l dNTPs, Buffer 10X, 25 mM MgCl₂, GC-Rich (10% of the final volume), 2 Units of Taq DNA polymerase (Applied Biosystems Inc., Branchburg, NJ), and 10 ng/µl of genomic DNA. Reactions were performed in a Perkin Elmer thermal cycler for one cycle at 94 °C for 6 min, 30 cycles at 94 °C for 40 s, 67 °C for 30 s, 72 °C for 45 s, and a final extension at 72 °C for 5 min. The amplified fragments, after digestion with 3 U of HhaI restriction enzyme, were separated using 5% agarose gel electrophoresis, and the restriction patterns were visualized by ethidium bromide staining and UV light. The genotypes of a group of patients were confirmed by ABI PRISM 310 Automated Sequencer (Applied Biosystems, Forster City, CA, USA).

Statistical analysis

Data are presented as mean value \pm SD. The population included in this study presented a normal distribution for age. Differences between groups were evaluated by one-way analysis of variance (ANOVA). When only two groups were present, unpaired t test was used. Correlation between variables was determined by linear regression analysis, while interactions between variables were calculated by correlation and multiple regression analyses. Covariate analysis was performed by z-test. All statistical procedures were performed using the StatView program (Abacus Concepts, Inc., SAS Institute, Cary, NC) [58].

Results

Descriptive analysis

The clinical characteristics of the total cohort and of the four subgroups (young SED, young ATHL, older SED and older ATHL) are reported in Table 1. The young and older cohort presented a mean age of 35.5 ± 9.6 and 60.4 ± 6.9 , respectively. The ATHL and SED groups did not present significant differences in body mass index (BMI) and age. As expected, the ATHL group presented a lower resting heart rate than the SED group (P<0.001). The level of physical activity was significantly higher in the ATHL group than the SED group in the total cohort as well as in young and older subjects.

When the cohort was stratified based on APOE e4 allele carriage (see below), there were 14 carriers of the allele (8 SED and 4 ATHL).

Expression of α -syn, tau, A β and of α -syn heterocomplexes with tau or A β in red blood cells: immunoblotting analysis

First, the presence of α -syn, tau and A β in RBCs was assessed by western blotting analysis. As depicted in Figure 1a, in RBCs, the anti- α -syn antibody recognized 15 kDa and 30 kDa proteins, corresponding to α -syn [61], whereas the anti-A β antibody labelled 5 and 15 kDa proteins (Fig. 1a), corresponding to A β monomeric and oligomer forms, respectively [62, 63]. Bands with a higher molecular weight than 25kDa (Fig. 1b, middle panel) could indicate elevated oligomeric form of A β [64, 65]. These data confirm that RBCs expressed detectable levels of α -syn and A β , consistent with data from the literature [45, 66]. Finally, the anti-tau antibody produced the characteristic triplet bands ranging between 55 and 74 kDa [67], together with an additional band, demonstrating that RBCs express the tau protein as well (Fig. 1a). Of note, bands lower than 50 kDa (Fig. 1b, lower panel have been related to truncated or cleaved forms of tau containing the C-terminal region [68].

Then, the presence of α -syn heterocomplexes in blood cells was verified using a coimmunoprecipitation-western blotting assay (Fig. 1b). To this purpose, cell lysates were immunoprecipitated using an anti- α -syn antibody and then immunoblotted with an anti-tau or anti-A β antibody. In parallel, lysates were also immunoprecipitated using an anti- α -syn antibody and immunoblotted with the same antibody. In α -syn immunoprecipitates obtained from RBCs (Fig. 1b, *upper panel*), the anti- α -syn antibody recognized 15 kDa and 30 kDa proteins that corresponded to α -syn protein [61]. The A β immunoblotting performed on α -syn immunoprecipitates from RBCs (Fig. 1b, *middle panel*) showed several immunoreactive bands that corresponded to the monomeric and oligomeric A β forms (Fig. 1b) [62]. Similar results were obtained probing α -syn immunoprecipitates with an anti-tau antibody (Fig. 1b, *bottom panel*). Globally, the results demonstrated that α -syn forms heterocomplexes with A β and tau at a peripheral level.

Establishment of an immunoenzymatic assay to detect a-syn/AB heterocomplexes

To confirm the qualitative results obtained in the co-immunoprecipitation-western blotting analysis and to quantitatively detect the levels of heterocomplexes in RBCs, an immunoenzymatic assay was developed and validated, as follows. Standard α -syn/A β heterocomplexes were prepared by incubating 1 mg of each protein, prepared in 2 mM SDS, in parafilm-sealed tubes at 37°C for 36 h with continuous mixing [14]. As depicted in Figure 2a (*left panel*), in α -syn immunoprecipitates, the anti- α -syn antibody recognized a 15 kDa protein corresponding to α -syn. In parallel, no significant labelling was detected in A β alone (Fig. 2a, *left panel*), confirming the specificity of the antibody immunoreactivity. The A β immunoblotting performed on α -syn immunoprecipitates revealed four immunoreactive bands (Fig. 2a, *right panel*) [62], demonstrating the induction of α -syn/A β heterocomplexes *in vitro* in our experimental conditions.

To set up the immunoenzymatic assay, eight different dilutions of α -syn/A β were prepared and captured on wells pre-coated with an antibody specific to A β ; the subsequent use of an antibody specific to α -syn and the appropriate HRP-conjugated antibody allowed for quantification of α -syn/A β (see Methods section for experimental details). The results (Fig. 2b) showed a concentration-dependent increase in the specific absorbance at 450 nm, thus demonstrating the specificity and validity of the assay. The absorbance at 450 nm of blank wells obtained in the absence of the α -syn primary antibody consistently remained under 20 % of the total values (data not shown).

To determine the amount of recombinant α -syn not bound to A β , different dilutions of recombinant α -syn/A β proteins were captured on wells pre-coated with an antibody specific to A β , as described above. Following incubation, the samples were collected, and α -syn levels (i.e., recombinant α -syn not bound to A β) were quantified by a validated immunoenzymatic assay. Such analyses revealed that in our experimental conditions (i.e., 2 mM SDS for 36 h), 85 ± 2 % of recombinant proteins formed α -syn/A β heterocomplexes (Fig. 2c). The corrected standard curve was then obtained by subtracting the amount of free α -syn from the theoretical α -syn/A β concentrations (Fig. 2c) and further used to quantify such heterocomplexes in blood cells.

Establishment of an immunoenzymatic assay to detect α-syn/tau heterocomplexes

Standard α -syn/tau complexes were prepared by incubating 1 mg of each protein, prepared in 2 mM SDS, in parafilm-sealed tubes at 37°C for 1 h. The tau immunoblotting performed on α -syn-immunoprecipitates revealed specific immunoreactive bands (Fig. 3a, *right panel, first line*) that were not shown for recombinant tau alone (Fig. 3a, *right panel, second line*). These data demonstrate the induction of α -syn/tau heterocomplexes *in vitro* in our experimental conditions.

Different dilutions of α -syn/tau were prepared and captured on wells pre-coated with a specific antibody to α -syn; the subsequent use of an antibody specific to tau and the appropriate HRP-conjugated antibody allowed for quantification of α -syn/tau. The results (Fig. 3b) showed a concentration-dependent increase in specific absorbance at 450 nm, thus demonstrating the specificity and validity of the assay.

To determine the amount of recombinant α -syn not bound to tau, different dilutions of recombinant α -syn/tau proteins were captured on wells pre-coated with a specific antibody to α -syn, as described above. Following incubation, the samples were collected, and tau levels (i.e., recombinant α -syn not bound to α -syn) were quantified by a validated immunoenzymatic assay. Such analyses revealed that in our experimental conditions (i.e., 2 mM SDS for 1 h), 78 ± 7 % of the recombinant proteins formed α -syn/tau heterocomplexes (Fig. 3c). The corrected standard curve was then obtained by subtracting the theoretical α -syn/tau concentrations from the amount of free α -syn (Fig. 3c) and further used to quantify such heterocomplexes in blood cells.

Total and oligomeric α-syn concentrations in RBCs of healthy subjects

Total and oligomeric α -syn levels were quantitatively measured in RBCs isolated from 106 healthy subjects (Table 2). In the total cohort of healthy subjects (young versus older), no correlations were found between age and total or oligomeric α -syn RBC concentrations (Fig. 4a and b, total α -syn: P=0,4123; oligomeric α -syn: P=0,8993). In contrast, total α -syn significantly decreased with age in the ATHL group (young ATHL versus older ATHL, Fig. 4a, P=0.0011), whereas an opposite trend was found in the SED group (young SED versus older SED, Fig. 4a, P=0.1376).

As depicted in Figure 4 (panel a and b), total and oligomeric α -syn in RBCs showed comparable values in the ATHL and SED groups, either in the total (total α -syn: P=0.6284, oligomeric α -syn: P=0.5012) or in the young population (young ATHL versus young SED, total α -syn: P=0.1683, oligomeric α -syn: P=0.9181). Moreover, in the older cohort, oligomeric α -syn was not significantly different between the ATHL and SED groups (Fig. 4b, P=0.1236). In contrast, total α -syn levels in

RBCs were significantly lower in the elderly ATHL group compared to that in the elderly SED group (Fig. 4a, P=0.0007), suggesting that physical activity may play a major role in modulating total α -syn levels in RBCs with increasing age.

Aβ concentrations in RBCs of healthy subjects

A β levels in RBCs (Table 2) were found to progressively accumulate with age in the SED population (young SED versus older SED, P=0.0369, Fig. 4c), whereas comparable values were found in the total (total young versus total older, P=0.5622) and ATHL population (young ATHL versus older ATHL, P=0.2502). Conversely, the analyzed parameter did not significantly differ between the ATHL and SED populations in the whole cohort (Fig. 4c, P=0.9407), as well as in the young subjects (young ATHL versus young SED, Fig. 4c, P=0.2987). Interestingly, as shown for total α -syn, the ATHL group presented significantly lower levels of A β in the older subgroup (older ATHL versus older SED, Fig. 4c, P=0.0360).

Tau concentrations in RBCs of healthy subjects

RBC levels of total tau significantly decreased with age in the total population (Fig. 4d, young versus older, P=0.0434); consistently, tau levels were significantly higher in the young SED population than in the older SED population (Fig. 4d, P=0.0202), while they did not change with age in the ATHL group (young ATHL versus older ATHL, P=0.5382).

Surprisingly, tau concentrations were significantly lower in the ATHL group than in the SED group in the total population (total ATHL versus total SED, Fig. 4d, P=0.0182) and in the young cohort (young ATHL versus young SED, P=0.0343) but not in the older one (older ATHL versus older SED, P=0.2893). These data suggest that RBC tau can exert beneficial effects especially in the young subgroup.

α -syn heterocomplexes with A β or tau in RBCs of healthy subjects

Finally, α -syn/A β and α -syn/tau heterocomplexes were measured in RBCs using the "home-made" immunoenzymatic assay. The results (Fig. 4e and f) showed that α -syn/A β or α -syn/tau levels in RBCs did not differ between young and elderly subjects (P=0.3529 and P=0.7145, respectively). Consistently, such parameters were comparable between the young and older SED groups and between the young and older ATHL groups (Fig. 4e and f).

Significantly lower concentrations of α -syn/A β were found in ATHL subjects than in SED subjects in the whole cohort (P=0.0001). This difference was also observed in the older subpopulation (older ATHL versus older SED, Fig. 4e, P=0.0055) and in the young one (young ATHL versus young SED, P=0.0038). These data suggest that α -syn/A β levels in RBCs are modulated by physical activity.

 α -syn/tau levels did not differ in young versus older subgroups (Fig. 4f and Table 2). Moreover, contrary to α -syn/A β , no significance difference in α -syn/tau concentrations (Fig. 4f) was found between the SED and ATHL groups (whole cohort: P=0.8659; older cohort: P=0.6021; young cohort: P=0.8993). These results suggest that the interaction of α -syn with tau is not modulated by physical exercise.

Plasma antioxidant capacity (AOC) in healthy subjects

The antioxidant capacity was measured in plasma from healthy subjects using the TOSC assay (Table 2); higher mean levels from this assay are related to a better antioxidant capability.

The results showed that AOC toward hydroxyl radicals was significantly higher in young subjects than in the elderly in the SED cohort (young SED vs older SED, P=0.0366, Fig. 5a), thus confirming that AOC progressively decreases with age [69].

No differences in TOSC values toward hydroxyl or peroxyl radicals were obtained when comparing young versus older subjects of the total population (Total young vs Total older, hydroxyl: P=0.2511; peroxyl: P=0.1640, Fig. 5a and b), and the young ATHL group showed significantly lower TOSC values toward hydroxyl than the older ATHL group (Fig. 5a, P=0.0019). These results suggest that additional factors other than age may influence AOC (see discussion section).

As expected, AOC toward hydroxyl (Fig. 5a) and peroxyl (Fig. 5b) radicals was significantly higher in the ATHL group than in the SED group in the whole cohort (Total ATHL vs Total SED, hydroxyl: P<0.0001; peroxyl: P=0.0442) and in the young subpopulation (Young ATHL versus young SED, hydroxyl: P=0.0362; peroxyl: P=0.0434, Fig. 5a and b). In the older subgroup (older ATHL versus older SED), a statistical significance was reached only for hydroxyl radicals (Fig. 5a and b, hydroxyl: P<0.0001; peroxyl: P=0.5360). These results confirm that physical activity is able to enhance antioxidant capacity in human subjects.

Consistent with these findings, hydroxyl TOSC values directly correlated with the level of physical activity in the whole subpopulation (Fig. 5c, P=0.0172, R²=0.131) and in older subjects (Fig. 5d, P=0.0055, R²=0.373) but not in young subjects (Fig. 5e, P=0.2352). These data indicate that physical activity level can influence AOC, in particular against hydroxyl radicals.

Interestingly, peroxyl TOSC values directly correlated with the physical activity level in the SED group (P=0.0188, R²=0.274, Fig. 5f), suggesting that a low grade of physical activity can play a role in the antioxidant capability against peroxyl radicals.

Correlation of ND-related proteins with plasma antioxidant capability

Correlations between plasma AOC and α -syn, A β , tau and α -syn heterocomplexes were determined by linear regression analysis.

1) α -syn: The total α -syn concentration in RBCs was inversely correlated with hydroxyl AOC in the whole cohort (Fig. 6a, P=0.0006, R²=0.251), as well as in the older subjects (Fig. 6b, P=0.0021, R²=0.435) but not in young subjects (Suppl. Fig. 1a, P=0.2656). Furthermore, hydroxyl TOSC values were inversely correlated with RBC total α -syn in the ATHL subgroup (Fig. 6c, P=0.0004, R²=0.409). Any significant correlation between hydroxyl TOSC and total α -syn concentration was found in SED group (P=0.223, R²=0.0097, Suppl. Fig. 1b). Based on these findings, a higher contribute of hydroxyl AOC on α -syn accumulation can be speculated with advanced age or in pre presence of physical exercise.

In contrast, no correlation was found between the concentration of oligomeric α -syn of RBCs and hydroxyl AOC in the whole cohort (Suppl. Fig. 1c, P=0.3874), the older subpopulation (Suppl. Fig. 1d, P=0.2961), or in the young one (Suppl. Fig. 1e, P=0.9458). These findings suggest that oligomeric α -syn is poorly related to hydroxyl AOC.

In contrast to hydroxyl AOC, peroxyl radicals did not show any correlation with total α -syn in the whole cohort (Suppl. Fig. 1f, P=0.4675), older subjects (Suppl. Fig. 1g, P=0.7630) or in the young ones (Suppl. Fig. 1h, P=0.5187).

2) A β : Peroxyl and hydroxyl TOSC values did not show any significant correlation with RBC A β levels in the total cohort (Suppl. Fig. 2a and b, peroxyl: P=0.2550; hydroxyl: P=0.197) or in young subjects (Suppl. Fig. 2c and d, peroxyl: P=0.2550; hydroxyl: P=0.197). Similar results were obtained in the SED and ATHL populations (Suppl. Fig. 2e-h, peroxyl SED: P=0.3837; hydroxyl SED: P=0.3435; peroxyl ATHL: P=0.7064; hydroxyl ATHL: P=0.7635). Interestingly, A β levels were inversely correlated with AOC toward hydroxyl radicals in the elderly cohort only (Fig. 6d, P=0.0342, R²=0.266), thus suggesting that A β levels are related to hydroxyl AOC with increasing age.

3) Tau: Peroxyl and hydroxyl TOSC values did not show any significant correlation with RBC tau levels, regardless of age (Suppl. Fig. 3a-f, total cohort, peroxyl: P=0.1563; hydroxyl: P=0.1563; young cohort: hydroxyl: P=0.5840; peroxyl: P=0.5528; older cohort, peroxyl: P=0.5943; hydroxyl:

P=0.1806), although a trend toward an inverse correlation was evidenced. Interestingly, tau levels in RBCs were inversely related to hydroxyl but not peroxyl TOSC values in the ATHL group (hydroxyl: P=0.0497, R^2 =0.151, Fig. 6e; peroxyl: P=0.5122, Suppl. Fig. 3g).

4) α -syn/A β : Hydroxyl TOSC values were inversely correlated with α -syn/A β levels in the whole population (Fig. 7a, P=0.0002, R²=0.287), older subjects (Fig. 7b, P=0.0239, R²=0.266), and the young subpopulation (Fig. 7c, P=0.0041, R²=0.318), indicating that a higher plasma AOC toward hydroxyl radicals is associated with a lower concentration of α -syn/A β heterocomplexes. Similar data were found in the ATHL subgroup (Fig. 7d, P=0.0003, R²=0.428), but not in SED (Fig. 7e, P= 1499). These data suggest that α -syn/A β accumulation is strongly associated with hydroxyl-related oxidative stress in RBCs.

No significant correlation between peroxyl TOSCA values and α -syn/A β levels was found, regardless of age (total population: P=0.5817; young: P=0.8817; older: P=0.4378) or physical activity (ATHL: P=0.0951; SED: P=0.1490).

5) α -syn/tau: An inverse correlation between α -syn/tau concentration and hydroxyl TOSC values was demonstrated in the whole population (Fig. 8a, P=0.0325, R²=0.107), the older subpopulation (Fig. 8b, P=0.0478, R²=0.196), and in the ATHL group (Fig. 8c, P=0.0478, R²=0.196). No relationship was detected in the SED group (Fig. 8d, P=0.1338) or in the young cohort (Fig. 8e, P=0.5548). These findings suggest that the interaction of α -syn with tau is modulated by hydroxyl AOC, particularly with increasing age and with a higher rate of physical exercise.

No correlation with TOSC against peroxyl radicals was detected, regardless of age (whole population: P=0.5968; older cohort: P=0.8385; young cohort: P=0.3640, Suppl. Fig. 4a-c), or physical activity (ATHL: P=0.9618; SED: P=0.3458, Suppl. Fig. 4d-e).

Correlation of ND-related proteins with the level of physical activity

Correlations between the level of physical activity and α -syn, A β , tau and α -syn heterocomplexes were determined by linear regression analysis.

1) α -syn: In RBCs of elderly subjects, both total and oligomeric α -syn were inversely correlated with physical activity level, although a statistical significance was only observed for total α -syn (Fig. 9a, total α -syn: P=0.0463, R²=0.103; Suppl. Fig. 5a, oligomeric α -syn: P=0.1256). In contrast, there was no significant correlation with the physical activity level in the total cohort (Suppl. Fig. 5b and c, total α -syn: P=0.6950; oligomeric α -syn: P=0.4842) or the young one (Suppl. Fig. 5d and e, total: P=0.5289; oligomeric: P=0.9529). Moreover, any significant correlation was found in the ATHL group (Suppl. Fig. 5f and g, total α -syn: P=0.8662; oligomeric α -syn P=0.8217). Interestingly, the grade of physical exercise was inversely related to oligomeric α -syn in the SED

group (Fig. 9b, P=0.0137, R²=0.099). These findings suggest that the rate of physical activity can partially modulate total α -syn accumulation, in particular with increasing age.

2) A β : A β concentrations in RBCs were directly related to the physical activity level in the young subgroup (Fig. 9c, P=0.0207, R²=0.083) and in ATHL (P=0.0473, R²=0.097, Fig. 9d). Any significant correlation was found in the whole population (P=0.4605, Suppl. Fig. 6a), in the elderly (P=0.1010, Suppl. Fig. 6b), or in SED (P=0.9268, Suppl. Fig. 6c). These data are consistent with the trend toward lower A β levels observed in the young ATHL group than in the young SED group (see Fig. 4c).

3) Tau: RBC tau concentrations were found to be inversely related to physical activity level in the total cohort (Fig. 9e, P=0.0284, R²=0.046). No other significant correlation was found in the other subgroups (Suppl. Fig. 6d-g, older: P=0.2397; young: P=0.9908; ATHL: P=0.9908; SED: P=0.5029).

4) α -syn/A β : The grade of physical exercise was inversely correlated with α -syn/A β concentrations of RBCs in the total population (Fig. 9f, P=0.0001, R²=0.134), as well as in young subjects (Fig. 9g, P=0.0006, R²=0.170) and in elderly (Fig. 9h, P=0.047, R²=0.108). These results suggest that the interaction of α -syn with A β is strongly modulated by the level of physical activity, independent of age. In contrast, no correlation was found in the ATHL (Suppl. Fig. 7a, P=0.3748) or SED populations (Suppl. Fig. 7b, P=0.8695).

5) α -syn/tau: No significant correlation between the physical activity level and α -syn/tau was detected (whole population: P=0.6326; older: P=0.8851; young: P=0.3915; ATHL: P=0.3915; SED: P=0.3574, Suppl. Fig. 7c-g), suggesting that the interaction of α -syn with tau is poorly modulated by the level of physical activity.

Covariate analysis

In a covariate analysis, as expected, total α -syn predicted the content of its oligomeric form (Z=2.969, P=0.0030). Interestingly, α -syn/A β complexes were shown to be a significant predictor for RBC concentrations of both total (Z=3.392, P=0.0007) and oligomeric (Z=3.136, P=0.0017) α -syn.

AOC toward hydroxyl radicals was demonstrated to be a significant predictor for total α -syn (Z=-3.485, P=0.0005) and α -syn/A β complexes (Z=-3.786, P=0.0002). Moreover, the physical activity level was shown to be a significant predictor for AOC toward hydroxyl radical (P=0.0166), tau (P=0.0282), and, especially, α -syn/A β complexes (P=0.0001).

Influence of the Apo E genotypes

Apo E, a plasma protein involved in lipoprotein metabolism [70], presents three major human isoforms, designated APO ε 2, APO ε 3, and APO ε 4. Among these, the APO ε 4 allele, leading to enhanced levels of brain A β , has been strongly associated with age-related diseases, including AD [71]. Furthermore, oxidative stress [70] and physical activity [42, 72] have been shown to differently act on the ε 4 population. On this basis, the present total population was stratified into ε 4-and non- ε 4 carriers (Table 1).

As depicted in Figure 10 (panel a and b), in the non ϵ 4-carriers, total α -syn (P=0.8544), oligomeric α -syn (P=0.2663), A β (P=0.4015), and α -syn/tau (P=0.1443) levels did not significantly differ between the ATHL and SED population, which is consistent with the data obtained in the whole population. Conversely, tau (Fig. 10b, P=0.0333) and α -syn/A β concentrations (Fig. 10b, P=0.0346) were confirmed to be lower in the ATHL than in the SED population. Moreover, the interaction of α -syn with A β was found to be inversely correlated with plasma AOC against hydroxyl radicals (Fig. 10c, P=0.0113, R²=0.195) and with the level of physical exercise (Fig. 10d, P=0.0292, R²=0.067). Similar to the results seen in the whole population (see Fig. 4), no significance differences between the SED and ATHL groups were found in RBC levels of total and oligomeric α -syn, A β , or α -syn/tau in ϵ 4 carriers (Fig. 11a and b). Interestingly, the concentration of α -syn/A β heterocomplexes (P=0.0421, Fig. 11b) were significantly lower in the ϵ 4-ATHL group than in the ϵ 4-SED group. Of note, α -syn/A β levels were inversely correlated with peroxyl TOSC values (Fig. 11c, P=0.0090, R²=0.982).

Unfortunately, the number of subjects carrying the ϵ 4 allele was too low to further divide into young and older cohorts.

Discussion

In the present study, the accumulation of misfolded proteins linked to NDs were related to the antioxidant capability (AOC) and physical activity in a cohort of 106 healthy subjects. The main conclusions of this work are as follows: i) the levels of tau and A β were differently modulated with increasing age; ii) total α -syn and A β accumulation in elderly subjects showed an inverse correlation with AOC toward hydroxyl radicals and the level of physical activity; iii) for the first time, α -syn was demonstrated to interact with A β and tau at a peripheral level; iv) α -syn heterocomplexes were strongly related to hydroxyl AOC; and v) α -syn/A β concentrations were inversely correlated with the level of physical activity.

These results suggested that α -syn heterocomplexes could be novel putative indicators to monitor antioxidant capacity and ND-related protein misfolding.

Oxidative stress has been suggested to be one of the potential common etiologies in various NDs because of its capability to trigger mitochondrial dysfunction, cellular damage, and an impairment of the DNA repair system, all of which have been known to be key factors in accelerating the aging process and ND development [1, 3, 28]. Particular attention has been paid to the relationship between oxidative stress and the accumulation of misfolded proteins, such as α -syn, A β and tau, which constitute the neuropathological hallmarks of AD, PD and other neurodegenerative proteinopathies [6]. For example, oxidative stress has been shown to exacerbate A β production and aggregation, as well as to promote tau phosphorylation, potentially inducing a vicious cycle of pathogenesis in AD [1, 73]. Similarly, mitochondrial dysfunction related to oxidative stress has been strongly associated with α -syn accumulation and apoptosis of dopaminergic neurons in PD [74].

Misfolded proteins related to NDs are hypothesized to accumulate in the brain even decades before the appearance of symptoms [75-77]; furthermore, recent studies have demonstrated a cell-to-cell transmission of pathologic α -syn and A β in anatomically interconnected areas [78, 79]. Brain, CSF and blood concentrations of such protein aggregates seem to be in a dynamic equilibrium [80, 81], suggesting that increased production in the brain could be associated with increased concentrations in the blood as the result of oligomer transfer across the blood brain barrier [82-84]. Among blood cells, RBCs have been suggested to be particularly sensitive to oxidative stress and misfolded proteins [45, 46, 85], exhibiting damage to cell membranes and decreased cell deformability, which is necessary for effective oxygen transport and delivery [85]. Based on these findings, recent efforts to study RBC concentrations of misfolded proteins and their relationship with oxidative stress or NDs have emerged [45, 46]. In this context, preliminary data have shown a correlation between A β concentrations in the brain and RBCs [42, 45], suggesting that these blood cells are a good model to study alterations in the brain. Herein, RBCs, isolated from a cohort of 106 healthy volunteers, were used to measure the accumulation of α -syn, A β , tau, and of their heterocomplexes, depending on the extent of the antioxidant capability or of physical exercise, which has become an important preventive and therapeutic tool in AD and PD [35, 86].

In our cohort, the antioxidant capability toward hydroxyl radicals decreased significantly with age in the total population and in the SED cohort, consistent with literature data reporting an inverse correlation between oxidative damage/antioxidant capacity and age [69, 87]. Surprisingly, AOC did not significantly differ between the young and elderly in the total cohort, and the young ATHL group showed significantly lower TOSC values toward hydroxyl than the older ATHL. We speculate that among the ATHL subgroup, additional factors, such as type and strength of exercise or the time since the last training, may interfere with such results. Consistent with this hypothesis, several factors, including diet, have been suggested to influence the extent of plasma AOC [44].

As expected, AOC toward hydroxyl and peroxyl radicals was significantly higher in the ATHL group than in the SED group, and the first directly correlated with the level of physical activity in the whole subpopulation and in older subjects. In this respect, regular exercise has been demonstrated to induce an adaptive response that involves the up-regulation of the enzymatic antioxidant system, culminating in a regulation of cellular redox state in brain cells [36, 37].

In contrast to plasma AOC, total and oligomeric α -syn did not significantly change with age in the whole population. Consistent with these data, no correlation between RBC α -syn oligomer levels and age has been found in PD patients [44]. Conversely, a recent paper has reported that plasma α -syn levels correlated strongly with age, revealing much lower concentrations in older than younger individuals [85]. This discrepancy may be explained by considering the different peripheral fluids under examination or the sensitivity of the antibodies in the home-made immuno-enzymatic assays.

Interestingly, total α -syn significantly decreased with age in the ATHL population, whereas an opposite trend was found in the SED group, suggesting that physical activity could differently modulate α -syn accumulation depending on age. Consistent with this hypothesis, total α -syn levels in RBCs were significantly lower in the ATHL group than in the SED group in the elderly (but not in the young or total) population and were inversely correlated with physical exercise level and hydroxyl AOC. Overall, these findings suggest that plasma AOC and the rate of physical activity can partially modulate total α -syn accumulation, in particular with increasing age. Consistent with our data, treadmill physical exercise has been shown to reduce α -syn and oxidative stress in a rat model of PD [88].

 $A\beta$ levels in RBCs were found to progressively accumulate with age in the SED population, consistent with previous data from the literature obtained in the same blood cells [46]. Nevertheless,

as shown for total α -syn, the influence of oxidative stress and physical activity on A β accumulation was relevant in the elderly subgroup only, in which A β levels were inversely correlated with AOC toward hydroxyl radicals and were significantly lower in the ATHL group than in the SED group. Consistent with our findings, RBC A β concentration positively correlates with an oxidative stress marker [46], and plasma A β concentrations decreased with physical exercise in elderly subjects (mean age 70) [42]. Moreover, a relationship between physical activity and brain amyloid load, quantified by PET, has been reported in a few studies [72, 89].

RBC levels of total tau significantly decreased with age in the total and SED populations; in contrast, Sparks and co-workers have reported a slight but significant age-related increase in circulating tau among individuals maintaining a cognitive control status [90]. This discrepancy can be explained considering that the authors analyzed tau plasma levels. Further studies will be needed to clarify the significance of tau in RBCs decrease with age.

The tau concentration was lower in the ATHL population than in the SED population and was inversely related to physical activity level in the total cohort, thus suggesting that this parameter is modulated by physical exercise. Consistent with this hypothesis, long-term physical exercise has been shown to reduce both total and hyperphosphorylated tau in transgenic mice [91].

Then, α -syn heteromeric association with A β or tau was analyzed in RBCs. α -syn and tau have been shown to co-localize in neurons [13, 17, 18] and within the axon compartment [19]; similarly, membrane-associated α -syn has been demonstrated to interact with A β [14, 15, 92, 93]. However, the presence of α -syn complexes with A β or tau in peripheral cells has not been investigated. Herein, by co-immunoprecipitation and immune-enzymatic assays, α -syn was shown to form heterocomplexes with tau and A β . The interaction of α -syn with tau or A β in RBCs did not change with age; conversely, both α -syn/A β and α -syn/tau concentrations were inversely correlated with hydroxyl AOC in the whole population and in older subjects. These data suggest that a higher plasma AOC toward hydroxyl radicals paralleled with lower concentrations of α -syn heterocomplexes.

The levels of α -syn/tau did not change with physical activity; conversely, significantly lower concentrations of α -syn/A β were found in the ATHL group versus the SED group in the whole cohort, as well as in the older and young subpopulation. In the three subgroups, the level of physical exercise inversely correlated with α -syn/A β concentrations of RBCs, suggesting that α -syn/A β levels in RBCs are modulated by physical activity, independent of age.

Finally, because only non ɛ4-carriers have been shown to receive the benefits of antioxidants [69] or physical activity [42], the influence of the Apo E genotype on the content of protein aggregates was analyzed. Tau levels remained significantly higher in the ATHL group than in the SED group,

suggesting that this parameter is not modulated by the Apo E isoform. As discussed above, the reduction in A β was only significant in older subjects; unfortunately, the low number of ϵ 4-carriers did not allow us to further divide the group into young and older subjects, which would have allowed for the evaluation of the influence of the Apo E genotype on A β accumulation. Conversely, despite the low number of subjects, α -syn/A β concentrations were inversely correlated with AOC against peroxyl radicals in the ϵ 4-population and were lower in the ϵ 4-ATHL subjects than in the ϵ 4-SED subjects. These data suggest that the interaction of α -syn with A β is not related to Apo E genotype, but rather, it can be influenced by the AOC toward peroxyl radicals in ϵ 4-carriers.

In conclusion, the negative role of oxidative stress and sedentary style in abnormal accumulation of ND-related proteins was shown in the present paper. Most importantly, for the first time, our data demonstrated the following: i) α -syn heterocomplexes with A β and tau were expressed in RBCs; ii) the interaction of α -syn with A β and tau was influenced by plasma AOC towards hydroxyl radicals; and iii) α -syn/A β content inversely correlated with the physical activity level. Together, all of the data suggest that α -syn heterocomplexes may represent potential new indicators to monitor antioxidant capacity and ND-related protein misfolding.

Future works will investigate the contribution of both oxidative stress and physical exercise on the accumulation of misfolded/hyperphosphorylated forms of ND-related proteins.

In interpreting such findings, it is important to note that our data were derived from a cohort of healthy subjects in peripheral fluids. Therefore, our results on α -syn heterocomplexes should be considered as preliminary data. Further research will be needed to establish a correlation between peripheral and central content of α -syn/A β and α -syn/tau, as well as to establish their role in pathological conditions of neurodegeneration. Finally, among the limitation of our study, the unfeasibility of measuring RBC turnover should be mentioned. Even if conflicting data on exercise-mediated modulation of RBC production have been reported [94-98], the influence of exercise on RBC ageing/turnover will be take into consideration in a future study.

Abbreviations:

AOC antioxidant capacity
Apo E Apolipoprotein E
Aβ β-amyloid₁₋₄₂
α-syn α-synuclein
AD Alzheimer's disease
DLB Dementia with Lewy Bodies
NDs Neurodegenerative diseases

PD Parkinson's disease
RBCs Red blood cells
ROS Reactive oxygen species.
TOSC Total Oxyradical Scavenging Capacity.

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Authors' contributions

SD, MLT, GS, UB and GS were involved with the conception, design, and interpretation of data. SD, DP, JF, CI, LP, ALG and LC performed the experiments. SD, DP, JF, FG, MLT, FF and FB were involved with data analysis. LP and FB collected the clinical material. CM, GS, MLT, GS and UB provided general overall supervision of the study, and acquired funding. All authors contributed to the drafting and critical revision of the manuscript and have given final approval of the version to be published.

Figure Legends

Fig. 1 Presence of α -syn, A β , tau and their heterocomplexes in blood cells. **a** Cell lysates obtained from RBCs were subjected to Western blot analysis using antibody to α -syn, A β or tau. GAPDH was the loading control. **b** Cell lysates obtained from RBCs were immunoprecipitated with an anti- α -syn antibody, and then immunoblotted with antibody to α -syn, A β or tau. One representative Western blot is presented for each condition.

Fig. 2 In vitro determination of α -syn/A β heterocomplexes. **a** Human recombinant α -syn/A β (first line) was immunoprecipitated with an anti- α -syn antibody, and then immunoblotted with antibody to α -syn or A β . Recombinant A β alone (second line) was used as a positive control. One representative Western blot is presented for each condition. **b**, **c** Different concentrations of human recombinant α -syn/A β were captured on wells pre-coated with an anti-A β antibody. After extensive washes, levels of the α -syn/A β complex were quantified using an antibody specific for α -syn, and subsequently an HRP-conjugated antibody and a TMB substrate kit. Blank wells were obtained in the absence of α -syn antibody. **c** The theoretical α -syn/A β concentrations were subtracted of the quote of free syn, not bound to A β . Data are expressed as absorbance at 450 nm minus blank values, and are the mean \pm SEM of at least three independent experiments.

Fig. 3 In vitro determination of α -syn/tau heterocomplexes. **a** Human recombinant α -syn/tau (first line) was immunoprecipitated with an anti- α -syn antibody, and then immunoblotted with antibody to α -syn or tau. Recombinant tau alone (second line) was used as a positive control. One representative Western blot is presented for each condition. **b**, **c** Different concentrations of human recombinant α -syn/tau were captured on wells pre-coated with an anti-tau antibody. After extensive washes, levels of the α -syn/tau complex were quantified using an antibody specific for α -syn, and subsequently an HRP-conjugated antibody and a TMB substrate kit. Blank wells were obtained in the absence of α -syn antibody. **c** The theoretical α -syn/tau concentrations were subtracted of the quote of free syn, not bound to tau. Data are expressed as absorbance at 450 nm minus blank values, and are the mean \pm SEM of at least three independent experiments.

Fig. 4 Determination of ND-related proteins in RBCs. **a-f** RBC levels of total α -syn (**a**), oligomeric α -syn (**b**), A β (**c**), tau (**d**), α -syn/A β (**e**) and α -syn/tau (**f**) in the total cohort, in young and older subgroups of ATHL and SED subjects (mean \pm SD). Lysates obtained from RBCs were subjected to specific immunoassay, as described in the Methods section. *P<0.05, ***P<0.001 versus other subgroups.

Fig. 5 Determination of plasma AOC in human subjects. **a**, **b** Plasma total oxyradical scavenging capacity (TOSC) against hydroxyl (**a**) and peroxyl (**b**) radicals in the total cohort, in young and older subgroups of ATHL and SED subjects (mean \pm SD). *P<0.05, **P<0.01, ***P<0.001 versus other subgroups. **c-f** Correlation analysis between TOSC values against hydroxyl or peroxyl radicals and level of physical activity, expressed as Physical Activity level.

Fig. 6 Correlation between ND-related proteins and plasma AOC. **a-c** Correlation analysis between total α -syn concentrations in RBCs (**a-c**) and TOSC values against hydroxyl radicals in the total cohort (**a**), older cohort (**b**), young cohort (**c**) or in ATHL (**d**). **e** Correlation analysis between A β concentrations in RBCs and TOSC values against hydroxyl radicals in the older cohort. **f** Correlation analysis between tau concentrations in RBCs and TOSC values against hydroxyl radicals in the older cohort. **f** Correlation analysis between tau concentrations in RBCs and TOSC values against hydroxyl radicals in ATHL.

Fig. 7 Correlation between α -syn/A β content in RBCs and plasma AOC. **a-d** Correlation analysis between α -syn/A β concentrations in RBCs and TOSC values against hydroxyl radicals in the total cohort (**a**), older cohort (**b**), young cohort (**c**), in ATHL (**d**) or in SED (**e**).

Fig. 8 Correlation between α -syn/tau content in RBCs and plasma AOC. **a-c** Correlation analysis between α -syn/tau concentrations in RBCs and TOSC values against hydroxyl radicals in the total cohort (**a**), older cohort (**b**), in ATHL (**c**), in SED (**d**) or in the young cohort (**e**).

Fig. 9 Correlation between ND-related proteins and physical exercise. **a-d** Correlation analysis between level of physical activity and RBC concentrations of total α -syn (**a**), oligomeric α -syn (**b**), A β (**c**), tau (**d**) and α -syn/A β (**e**, **f**, **g**) in the indicated subgroups.

Fig. 10 ND-related proteins in non ε 4-carriers. **a**, **b** RBC levels of total α -syn, oligomeric α -syn, A β , α -syn/A β , tau and α -syn/tau in subjects not carrying the ε 4 allele (total cohort, ATHL and SED subjects). Lysates obtained from RBCs were subjected to specific immunoassay, as described in the Methods section. The data are the mean \pm SD. *P<0.05 ATHL versus SED. **c**, **d** Correlation analysis between RBC concentrations of α -syn/A β and TOSC values against hydroxyl radicals (**c**) or level of physical activity (**d**) in subjects not carrying the ε 4 allele.

Fig. 11 ND-related proteins in ε 4-carriers. **a**, **b** RBC levels of total α -syn, oligomeric α -syn, A β , α -syn/A β , tau and α -syn/tau in subjects carrying the e4 allele (total cohort, ATHL and SED subjects). Lysates obtained from RBCs were subjected to specific immunoassay, as described in the Methods section. The data are the mean \pm SD. **P<0.01 ATHL versus SED. **c** Correlation analysis between

RBC concentrations of α -syn/A β and TOSC values against peroxyl radicals in subjects carrying the ϵ 4 allele.

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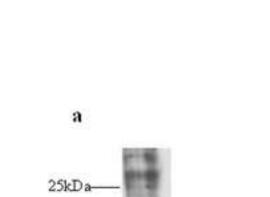
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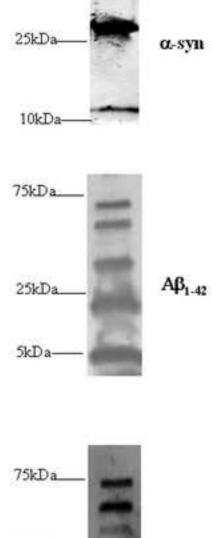
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Figure 1

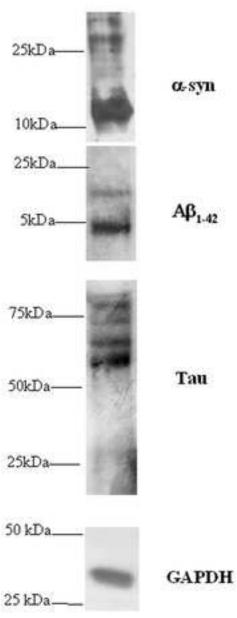


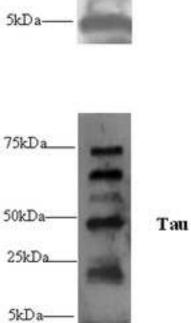


I.P. α- syn

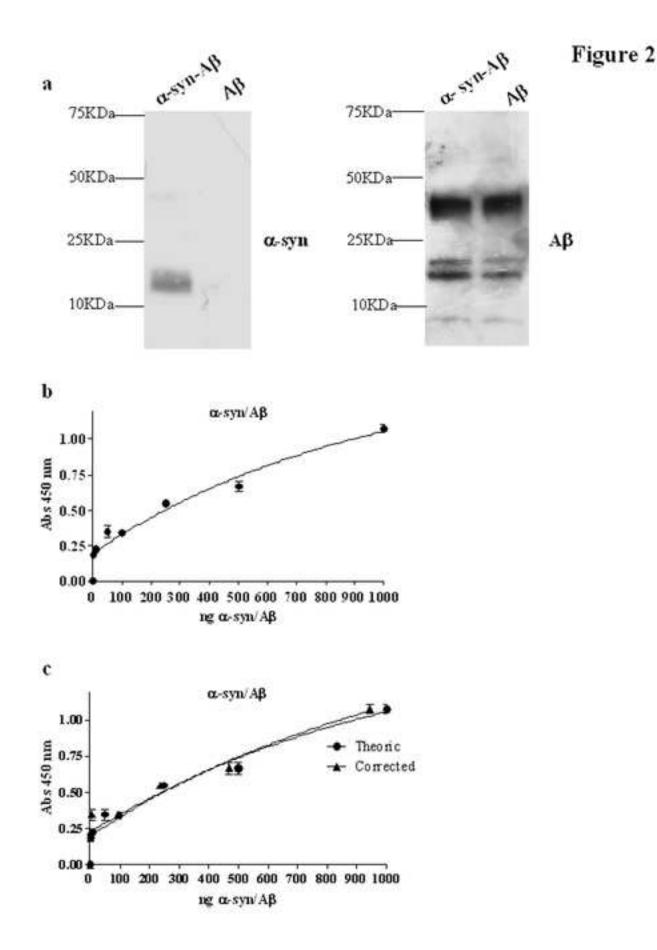
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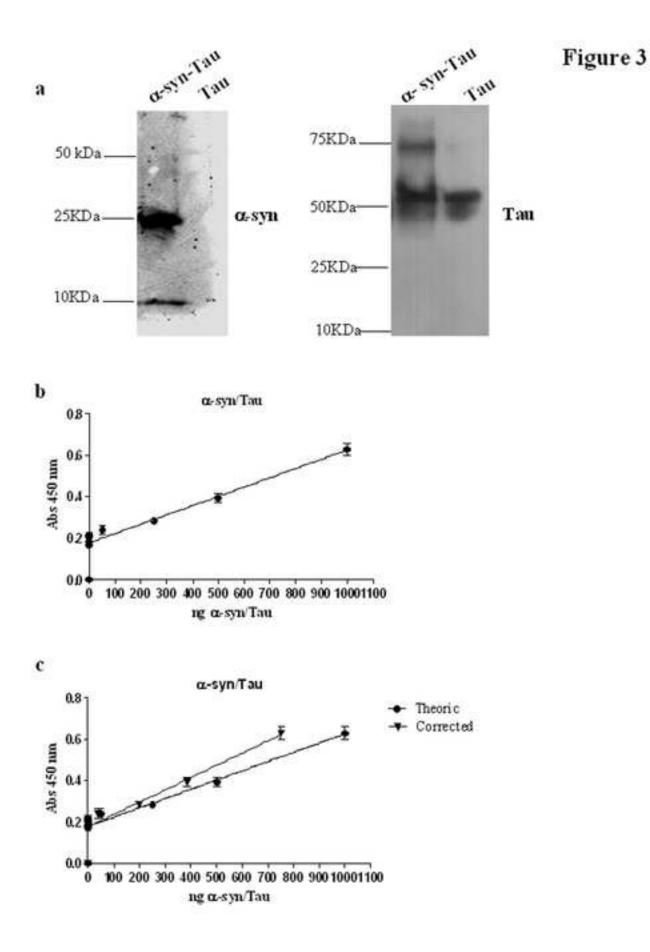


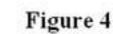


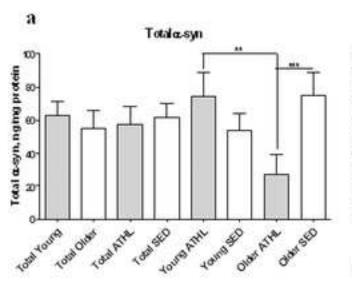


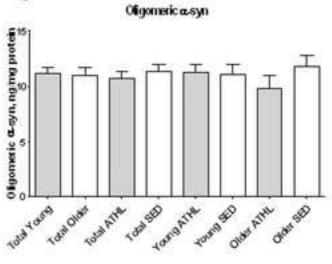






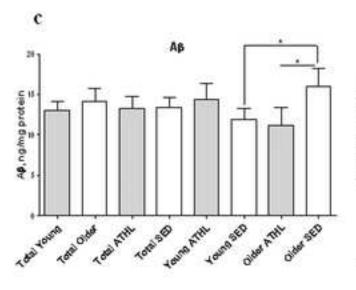


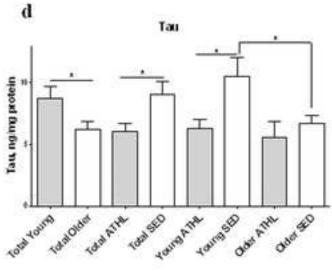




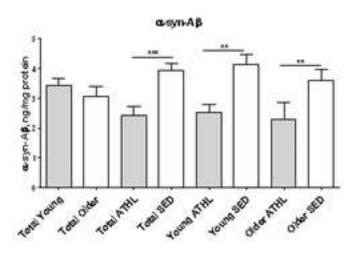
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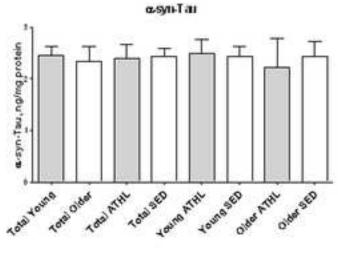
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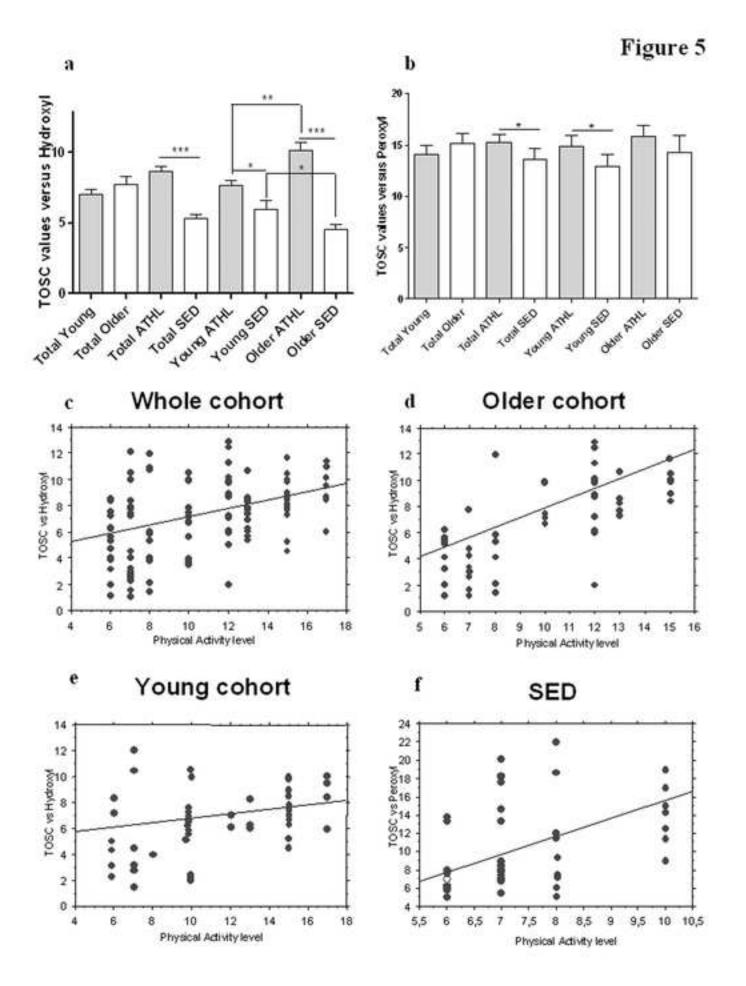




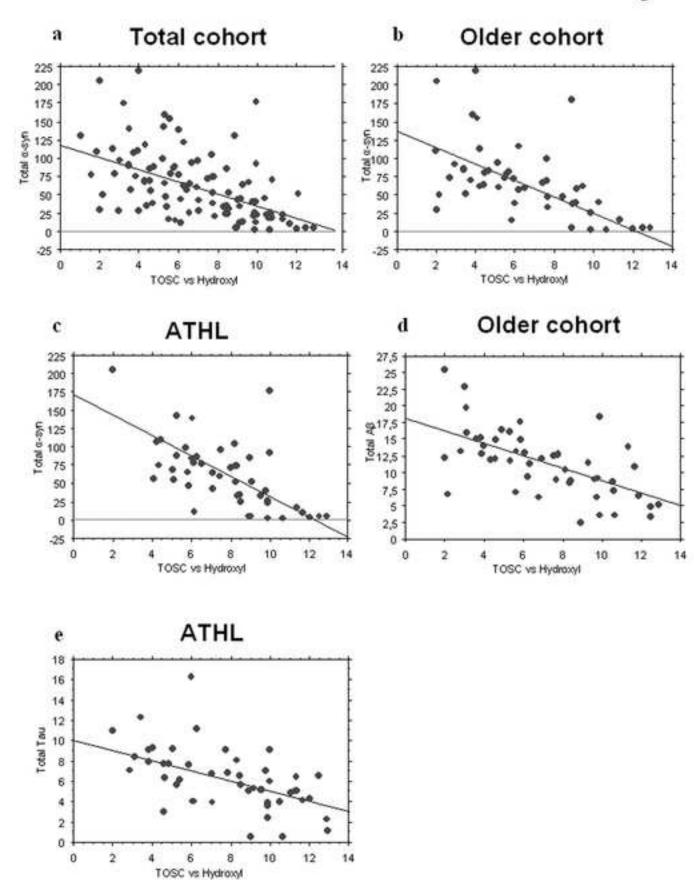












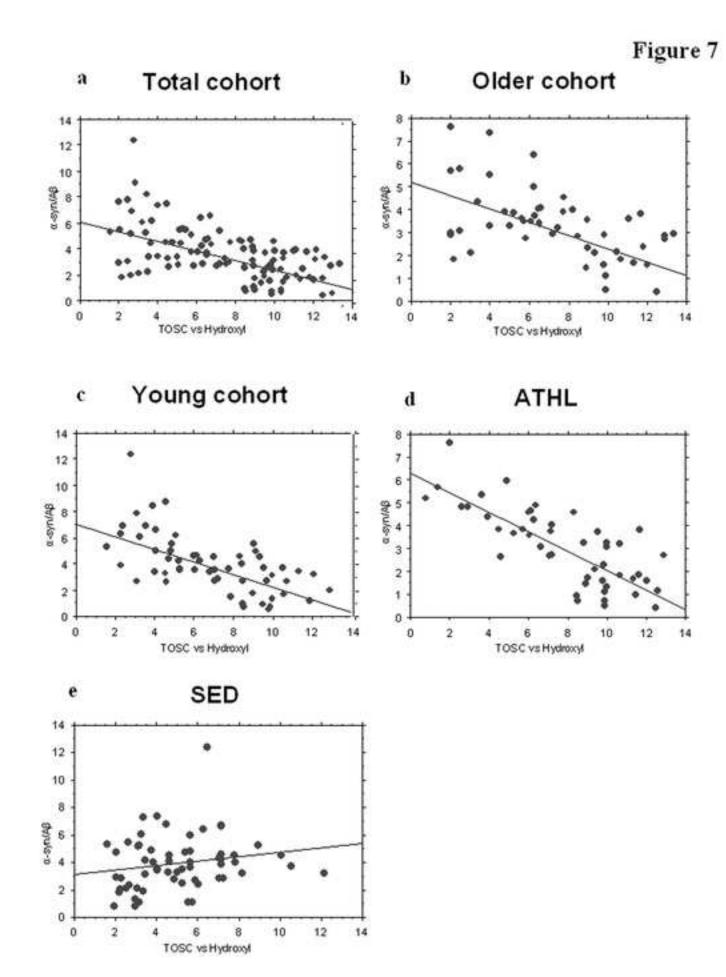
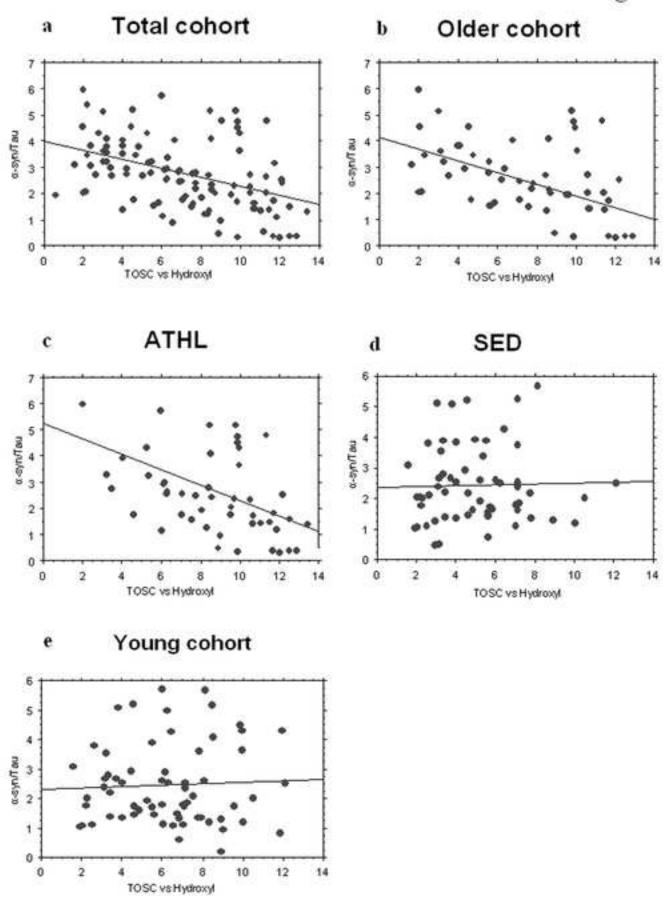
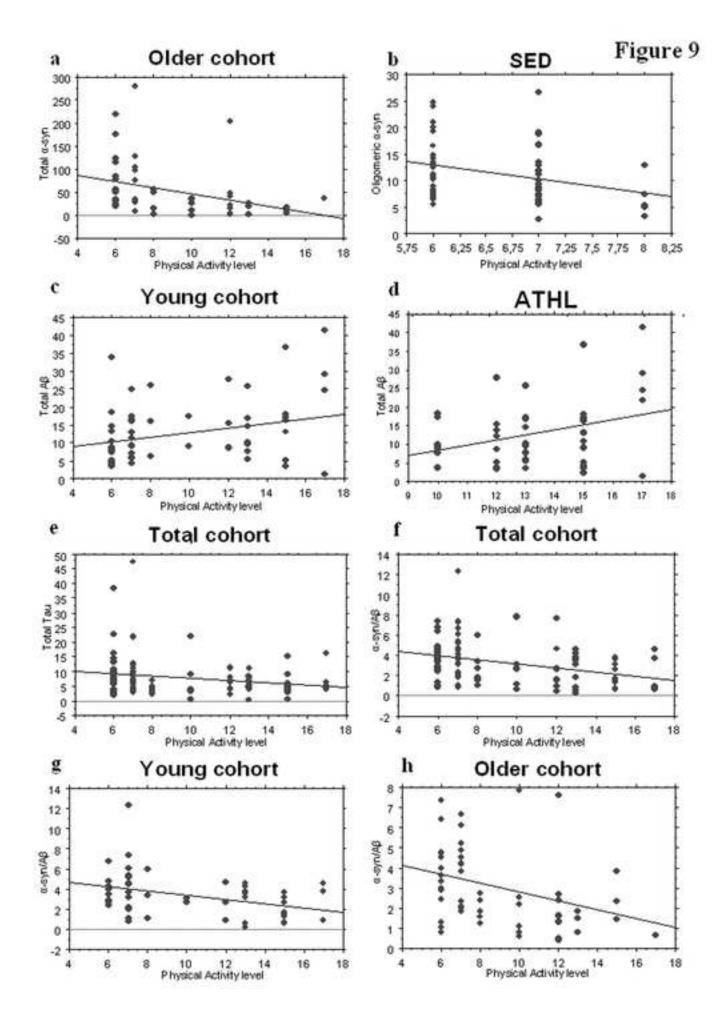




Figure 8

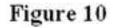


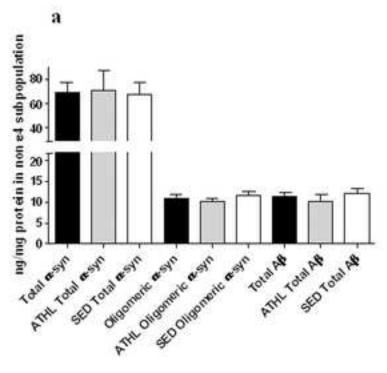


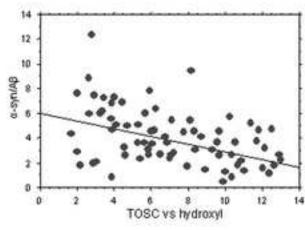
Non £4 carriers

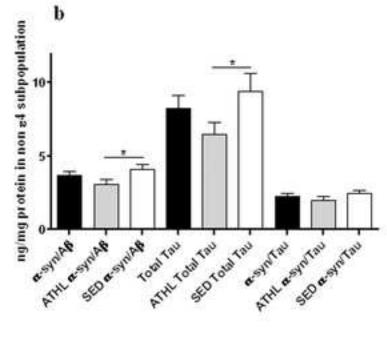
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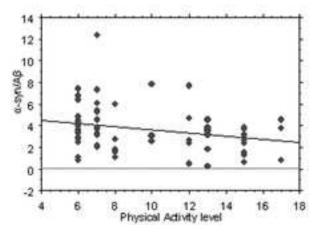
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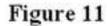












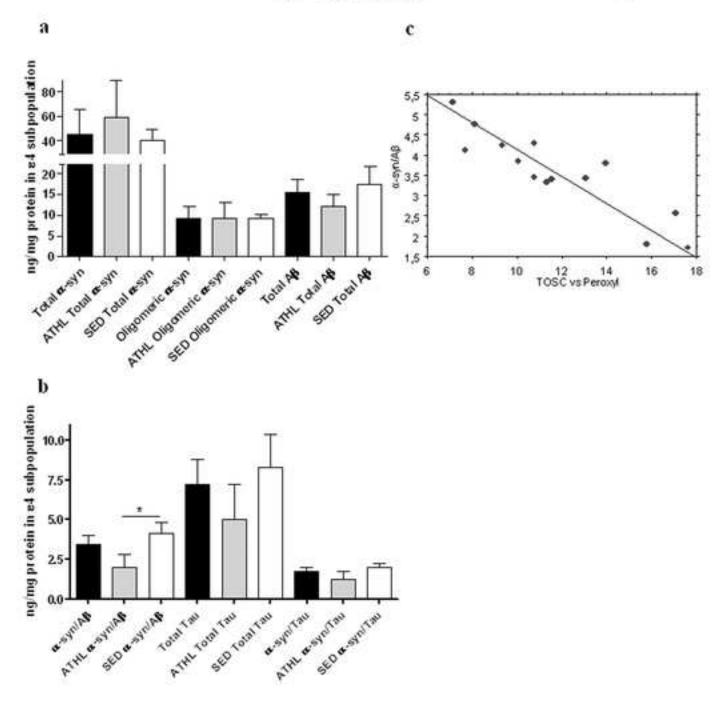


Table 1. Descriptive analysis of the total population and of the subgroups. The data are the mean±SD.

	Number of	Age (y)	BMI	Heart rate	Physical	
	subjects (N)				activity level	
Young subjects	56	35.5±9.6	23.8±1.8	55.4±3.3	9.71±3.83	
Older subjects	50	60.4±6.9	24.6±2.1	60.30±5.65	8.79±3.15	
ATHL (Total	48	44.6±13.5	23.8±1.8	52.6±3.6	13.2±2.2	
cohort)						
SED (Total cohort)	58	46.7±14.5	24.4±1.9	63.1±5.4	6.61±0.64	
Young ATHL	22	35.8 ± 8.0	23.4±1.6	50.2±3.9*	13.75±2.05	
Young SED	26	36.3 ± 9.2	24.2±1.20	60.5±2.6	6.70 ±0.59	
Older ATHL	22	57.4± 6.7	24.1±0.8	54.9± 3.2*	12.3±2.1	
Older SED	32	61.2±8.0	25.1±0.9	65.7± 8.1	6.50±0.68	
Non ɛ4 carrier	73	44.4±14.2	24.1±1.1	58.8±1.1	9.1±3.4	
ATHL non ε4	29	41.1±12.8	23.5±0.9	53.4±0.7	13.3±2.3	
carrier						
SED non ɛ4 carrier	44	46.6±14.7	24.6±1.2	64.2±1.4	6.57±0.66	
ε4 carrier	14	39.3±14.3	24.0±1.1	56.1±1.4	9,.0±3,.1	
ATHL ε4 carrier	6	41.5±21.0	23,7±0.8	51.2±1.2	13.3±2.6	
SED ɛ4 carrier	8	38.3±11.3	24.3±1.3	60.9±1.5	6.87±064	

BMI, Body Mass Index; ATHL, Athletes, SED, sedentary. * P < .001 vs sedentary subgroups

	TOSC values	TOSC values	Total α-syn	Oligomeric α-syn	Αβ	α-syn/Aβ	Tau	α-syn/tau
	(Peroxyl)	(Hydroxyl)						
Young subjects	14.1±3.5	7.05±2.74	62.6±50.0	11.1±5.0	13.0±9.2	3.44±1.99	8.70±7.68	2.46±1.34
Older subjects	15.2±4.3	7.79±3.82	54.9±45.9	11.0±4.7	14.1±10.2	3.07±2.09	6.21±4.11	2.35±1.74
ATHL (Total	15.3±3.9	8.70±2.65	57.4±43.8	10.7±4.3	13.3±9.7	2.45±1.79	6.05±4.20	2.39±1.79
cohort)								
SED (Total cohort)	14.0±2.7	5.34±2.05	61.6±44.6	11.3±5.3	13.4±9.6	3.94±1.97	9.07±7.83	2.44±1.24
Young ATHL	14.9±3.6	7.66±1.85	74.3±60.2	11.2±4.1	14.4±10.1	2.53±1.47	6.32±3.67	2.49±1.53
Young SED	12.9±3.0	6.03±3.10	53.8±40.8	11.0±5.7	11.9±6.5	4.14±2.07	10.5±8.3	2.44±1.19
Older ATHL	15.8±3.5	10.1±3.0	26.8±19.5	9.8±4.6	11.1±8.6	2.30±1.31	5.59±4.12	2.22±1.74
Older SED	15.3±3.0	4.57±2.10	74.5±59.8	11.8±4.7	16.0±7.9	3.60±1.80	6.67±3.28	2.44±1.34
Non ɛ4 carrier	15.4±4.3	7.06±3.24	68.6±53.3	11.1±5.3	11.3±8.4	3.69±2.03	8.26±7.55	2.27±1.43
ATHL non E4	16.8±3.1	8.28±2.97	70.6±66.0	10.2±4.4	10.3±7.8	3.07±1.83	6.45±3.92	1.96±1.60
carrier								
SED non ɛ4	13.7±4.9	5.69±3.06	67.3±64.7	11.6±5.9	12.0±8.8	4.09 ± 2.07	9.45±6.72	2.47±1.30
carrier								
ε4 carrier	11.4±4.5	5.19±3.18	45.7±20.3	9.18±3.03	15.5±11.0	3.45±1.99	7.23±5.49	1.76±0.89
ATHL ɛ4 carrier	13.5±5.9	7.63±1.91	59.0±30.4	9.4±3.96	12.1±6.1	2.03±1.55	5.02±4.47	1.27±0.99
SED ɛ4 carrier	9.3±3.1	2.76±1.73	40.3±25.1	9.20±2.78	17.3±12.7	4.17±1.86	8.53±5.89	2.00±0.79

Table 2, TOSC values against hydroxyl and peroxyl radicals; concentrations of total α -syn, oligemeric α -syn, A β , α -syn/A β , tau and α -syn/tau (expressed as ng/mg protein) in the indicate subgroups. The values are expressed as mean \pm SD.

BMI, Body Mass Index; ATHL, Athletes, SED, sedentary

Supplementary Material

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