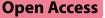
PERSPECTIVE



Cerebral apolipoprotein E and amyloid precursor-like protein 1 as risk factors for chronic neurodegeneration after non-traumatic acute brain injury (ABI)

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Acute brain injury (ABI) is defined as damage to the brain parenchyma as a result of an acute traumatic or non-traumatic (e.g., subarachnoid hemorrhage [SAH], ischemic stroke, intracerebral hemorrhage [ICH], acute demyelinated encephalopathy [ADE], acute hydrocephalus) injury [1, 2]. Patients with traumatic brain injury (TBI) are at an increased risk of developing chronic neurodegenerative diseases, including Alzheimer's disease (AD) [2], an association thought to be related to several factors, including altered production and clearance of apolipoprotein E (ApoE) and amyloid- β (A β) [3, 4]. ApoE is involved in lipid transport and injury repair in the brain. Lipidated ApoE binds to soluble A β and facilitates A β uptake through cell surface receptors, including

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⁴ Laboratory for the Structure and Function of Biological Membranes, Center for Structural Biology and Bioinformatics, Université Libre de Bruxelles, Brussels, Belgium low-density lipoprotein receptor (LDLR), LDLR-related protein 1 (LRP1), and heparan sulphate proteoglycan [4]. A β is produced by the cleavage of amyloid precursor proteins (APPs) in the cell membrane by b-secretase and g-secretase into a monomeric form with peptides of different lengths such as amyloid beta 1–40 (A β_{1-40}), and amyloid beta 1–42 (A β_{1-42}), which is then transformed into oligomeric and fibril forms and is one of the hallmarks of AD [5]. $A\beta_{1-40}$ and $A\beta_{1-42}$ speed up the aggregation kinetics and alter the pattern of spontaneously formed oligomeric species, which are considered the main toxic species [5-8]. Mounting evidence from genetic, pathological, and functional studies has shown that accumulation and aggregation of $A\beta$ in the brain is the result of an imbalance between its production and clearance. The A β peptides aggregate as toxic soluble A β oligomers, which then form amyloid plaques that injure synapses and ultimately cause neurodegeneration and dementia. After TBI, the concentration of ApoE in the cerebrospinal fluid (CSF) is low, which represents a risk factor for the development of Alzheimer's diseaselike disease [9, 10]. Additionally, reduced levels of CSF $A\beta_{1-42}$ accumulation in A β plaques, which are the pathological hallmark of AD, can be found in terminal axons within hours of death in up to 30% of patients with TBI [3]. However, data on the associations of ApoE and A β in patients with non-traumatic ABI have rarely been reported.



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We previously reported decreased CSF expression of proteins related to cholesterol metabolism in patients with ABI [1], which may potentially lead to reduced neuro-steroid production, increased risk of neurodegenerative disease, and worse functional outcomes. In a post hoc analysis of that study, we tested the hypothesis that low ventricular CSF (vCSF) concentrations of ApoE and A\beta-related proteins would also be present early after non-traumatic ABI and may be associated with unfavorable neurological outcome, defined as a Glasgow Outcome Scale score of 1-3 at 3 months after admission. In this post hoc analysis, ApoE (ApoE_{elisa}), amyloid beta 1–40 (A β_{1-40}), and amyloid beta 1–42 $(A\beta_{1-42})$ concentrations were measured in vCSF taken from an external ventricular drain on Days 1-5 after non-traumatic ABI. Control patients had scheduled elective clipping of a non-ruptured aneurysm and CSF was sampled just prior to surgery (Day 1). All samples were volume normalized. The vCSF concentrations of ApoE were measured using a commercial enzymelinked immunosorbent assay (ELISA) test (Quantikine® ELISA human apolipoprotein E/ApoE Immunoassay, R&D Systems® Minneapolis, MN) according to the manufacturer's protocol. The vCSF concentrations of $A\beta_{1-40}$ and $A\beta_{1-42}$ were also measured using a commercial ELISA test (human Aβ40 and Aβ42 ELISA kits, Invitrogen, Thermofisher USA) according to the manufacturer's protocol, with 50 μ l/patient (robotic pipette) per sample in duplicate using $A\beta_{1-40}$ and $A\beta_{1-42}$ monoclonal antibody (Biognost®, Germany), respectively. ApoE_{elisa} data were available for 19 patients with nontraumatic ABI (10 SAH, 2 ischemic stroke, 5 ICH, 1 ADE, 1 acute hydrocephalus) and for 5 control patients; data for $A\beta_{1-40}$ were available for 20 patients with non-traumatic ABI (11 SAH, 2 ischemic stroke, 4 ICH, 1 ADE, 2 acute hydrocephalus) and 10 controls; and data for $A\beta_{1-42}$ were available for 23 patients with nontraumatic ABI (SAH, n = 14; IS, n = 2; ICH, n = 5, ADE, n=1; acute hydrocephaly, n=1) and 10 controls. We also used data independent acquisition and SWATH mass spectrometry (SWATH-DIA) to measure ApoE (ApoE_{swath}), APP, and amyloid precursor-like 1 (APLP1) and 2 (APLP2) as previously described [1]; data using this technique were available from 29 patients with non-traumatic ABI (19 SAH, 3 ischemic stroke, 6 ICH, 1 acute hydrocephalus) and 10 controls. ELISA samples were taken only on Day 1 and SWATH-DIA on days 1-5. We used a Shapiro-Wilk's test to assess distribution and a non-parametric Wilcoxon test to quantify differences in vCSF proteomic expression between the groups. All tests were done using the latest version of R language [11]. The original study [1] was approved by the Erasme Hospital (Brussels, Belgium) Ethics Committee (2014/170-2015/130) and all patients or their next-of-kin signed written informed consent.

Day 1 ApoE_{elisa} concentrations were significantly lower (p < 0.01) in patients with non-traumatic ABI than in control patients. vCSF A β_{1-40} (p < 0.001) and A β_{1-42} (p < 0.001) concentrations were also lower in the ABI patients (Fig. 1). Day 1–5 ApoE_{swath} protein expressions were significantly lower in patients with nontraumatic ABI than in controls (p < 0.0001) (Additional file 1: Table S1). APLP1 values, but not those of APP or APLP2, were also significantly lower in non-traumatic ABI (p < 0.0001) throughout the 5-day period (Table S1). There were no differences in ApoE or A β proteins in patients with unfavorable and favorable neurological outcomes (data not shown).

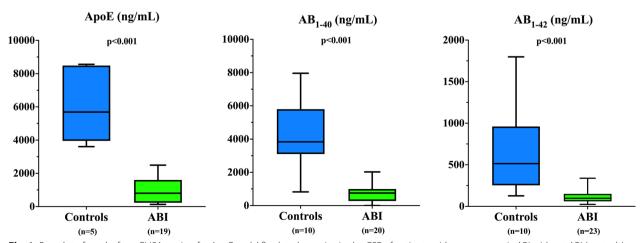


Fig. 1 Box plot of results from ELISA testing for ApoE and Aβ-related proteins in the CSF of patients with non-traumatic ABI without ABI (controls). Boxes represent the 25th–75th interquartile ranges with medians (horizontal line); the whiskers represent the minimum and maximum values in the dataset

These findings suggest that specific precursors of neurotoxic A β may be expressed in the early phase of nontraumatic ABI, such as that caused by vascular injury after ischemic or hemorrhagic strokes. However, this proteomic pattern in the vCSF was not associated with 90-day neurological outcome, which may be explained by various reasons. First, ABI cannot be entirely responsible for 90-day outcome, as many other factors contribute to functional recovery, such as central nervous system infection, seizures, rehabilitation, and pre-existing brain dysfunction. Second, the GOS is a measure of functional outcome rather than neurocognitive outcome and may not be the most suitable measure for assessing correlations between proteomic profiles and risk of developing neurocognitive disease. Third, not all the analyzed vCSF samples gave an ELISA result, which may have introduced a bias in the results. Lastly, the sample size was too small to allow normalization for age or comorbidities [1].

We acknowledge that this study is more valuable as an exploration of the proteomic profile after non-traumatic ABI rather than to assess its impact on neurological outcome. More studies on the vCSF proteomic profile of patients with ABI are needed to help better understand the cerebral response to an acute injury and the effect of such profiles on neurological outcomes in this setting.

Supplementary Information

The online version contains supplementary material available at https://doi.org/10.1186/s13054-023-04538-y.

Additional file 1. Supplemental Table S1. Comparisons of MS/MS swath results between patients without ABI (controls) and those with non-traumatic ABI. ApoE: Apolipoprotein E; ABI:40 amyloid beta 1:40; ABI:42 amyloid beta 1:42; APP: amyloid precursor protein; APLP1: amyloid precursor-like protein 1; APLP2: amyloid precursor-like protein 2. **ompared to Day 1 results for controls. **: MS/MS SWATH-DIA is a measure of abundance with no specific unit of quantification.

Acknowledgements

None.

Author contributions

CAS, DC and FT designed the study; CAS wrote the first draft of the manuscript; FT, JLV, DC, VR, VI, SB, JC reviewed the article for critical content; and all authors read and approved the final text.

Funding

None.

Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Declarations

Ethics approval and consent to participate

The original study from which the cohort derived was approved by the Erasme Hospital (Brussels, Belgium) Ethics Committee (2014/170-2015/130) and all patients or their next-of-kin signed written informed consent.

Consent for publication

Not applicable.

Competing interests

JLV is Editor-in-Chief of Critical Care. He has no other conflicts of interest. The other authors have no conflicts of interest to declare.

Received: 8 May 2023 Accepted: 20 June 2023 Published online: 24 June 2023

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