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Multiple Proteins Implicated in Neurodegenerative Diseases Accumulate in Axons After Brain Trauma in Humans

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Abstract

Studies in animal models have shown that traumatic brain injury (TBI) induces the rapid accumulation of many of the same key proteins that form pathologic aggregates in neurodegenerative diseases. Here, we examined whether this rapid process also occurs in humans after TBI. Brain tissue from 18 cases who died after TBI and from 6 control cases was examined using immunohistochemistry. Following TBI, widespread axonal injury was persistently identified by the accumulation of neurofilament protein and amyloid precursor protein (APP) in axonal bulbs and varicosities. Axonal APP was found to co-accumulate with its cleavage enzymes, beta-site APP cleaving enzyme (BACE), presentiin-1 (PS1) and their product, amyloid- β (A β). In addition, extensive accumulation of α -synuclein (α -syn) was found in swollen axons and tau protein was found to accumulate in both axons and neuronal cell bodies. These data show rapid axonal accumulation of proteins implicated in neurodegenerative diseases including Alzheimer's disease and the synucleinopathies. The cause of axonal pathology can be attributed to disruption of axons due to trauma, or as a secondary effect of raised intracranial pressure or hypoxia. Such axonal pathology in humans may provide a unique environment whereby co-accumulation of APP, BACE, and PS1 leads to intra-axonal production of A β as well as accumulation of α -syn and tau. This process may have important implications for survivors of TBI who have been shown to be at greater risk of developing neurodegenerative diseases.

Keywords

Traumatic brain injury; TBI; axonal injury; amyloid β; APP; BACE; PS-1; α-synuclein; tau

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Introduction

It has become increasingly accepted that traumatic brain injury (TBI) results in pathophysiological changes similar to those seen in neurodegenerative diseases. Several investigations have suggested a link between a history of TBI and the subsequent development of Alzheimer's disease (AD) (Mortimer et al. 1985; Rasmusson et al. 1995; Schofield et al. 1997; Nemetz et al. 1999; Guo et al. 2000; Lye and Shores 2000; Plassman et al. 2000). Likewise, TBI is an epidemiological risk factor for the development of sporadic Parkinson's disease (PD) (Nayernouri 1985; Factor and Weiner 1991; Stern 1991; Ben-Shlomo 1997; Lees 1997; Goldman et al. 2006).

Pathologically, AD is characterized by A β -containing plaques and neurofibrillary tangles comprised of tau protein (Braak and Braak 1991; Selkoe 2001; Forman et al. 2004). To a lesser extent, both dystrophic neurites and Lewy bodies containing α -synuclein protein (α syn) are also observed in AD. Lewy bodies and α -syn immunoreactivity are also hallmark pathological features of PD and other synucleinopathies such as dementia with Lewy bodies (DLB) and multi-system atrophy (MSA) (Smith et al. 2003; Norris et al. 2004). As with neurodegenerative diseases, protein accumulation is also a feature of TBI. Most notably, A β plaque formation and the accumulation of neurofilament proteins, tau and α -syn have been found in brain tissue of humans within hours to days following TBI (Grady et al. 1993; Roberts et al. 1994; Graham et al. 1995; Newell et al. 1999; Smith et al. 2003; Smith et al. 2003; Abrahamson et al. 2006). The mechanism underlying this rapid protein build-up after TBI remains unknown, as does its contribution to the later development of neurodegenerative disease.

Aß peptide is generated via the trans-membrane cleavage of amyloid precursor protein (APP) by the β - and γ -secretases. More specifically, its anabolism is mediated by beta-site APP cleaving enzyme (BACE) and the catalytic component of β -secretase, presenilin-1 (PS1) (De Strooper et al. 1998; Vassar et al. 1999; Nunan and Small 2000; Selkoe and Wolfe 2000; Esler and Wolfe 2001). Mounting evidence suggests that this process may also occur within the axonal membrane compartment. Large accumulations of A β have been found in swollen axons after TBI in a pig model of head rotational acceleration (Smith et al. 1999; Chen et al. 2004), in rodent models of brain contusion (Iwata et al. 2002; Stone et al. 2002; Chen et al. 2004), and in humans (Roberts et al. 1994; Smith et al. 2003). Axonal accumulations of A β were frequently found near diffuse, extracellular AD-like A β plaques in both the pig and in humans at the earliest survival timepoints measured (3 days and 18 hours respectively). This suggests a potential link between axonal pathology and A β plaque formation. (Smith et al. 1999, 2003b). More recently, extensive co-accumulations of A β with APP, BACE, and PS-1 were identified at sites of axonal injury and disconnection after TBI in the pig (Chen et al. 2004). Thus, disruption of axonal transport after TBI may create an environment whereby large accumulations of APP are processed to form $A\beta$, potentially leading to subsequent neurodegneration. Indeed, other recent studies have demonstrated the intra-axonal generation of A β in both central and peripheral nerve axons (Kamal et al. 2000; Kamal et al. 2001). Similarly, in a transgenic mouse model of AD, interrupted axonal transport and axonal swelling was shown to promote A β generation (Stokin et al. 2005).

The other classic pathological findings in AD are neurofibrillary tangles (NFTs) and neuropil threads (Braak and Braak 1991; Selkoe 2001; Forman et al. 2004). These intracellular structures are found to contain abnormal forms of the microtubule associated protein tau. NFTs with similarly abnormal tau are found in the brains of patients with dementia pugilistica; a progressive dementing disorder resulting from repetitive head trauma (Schmidt et al. 2001). Following a single episode of TBI in humans, hyperphosphorolyated tau has been demonstrated in brain tissue as well as elevated levels of the protein in cerebrospinal fluid (Newman SJ 1995; Zemlan et al. 1999). Additionally, excessive tau protein accumulation has been found in swollen axons in a pig model of TBI (Smith et al. 1999).

 α -syn is a small, highly soluble protein believed to play a role in synaptic maintenance (Norris et al. 2004). In the context of AD and the synucleinopathies, this protein is found as abnormal, highly insoluble, filamentous perikaryal aggregates (Trojanowski and Lee 2002; Forman et al. 2004). It appears that the α -syn found in disease states is pathologically altered due to ubiquitination, oxidation/nitration, phosphorylation and/or conformational modification (Giasson et al. 2000; Duda et al. 2002; Fujiwara et al. 2002). α -syn accumulation has also been demonstrated in neurons and axons following a single episode TBI in humans (Newell et al. 1999; Ikonomovic et al. 2004) as well as in patients with dementia pugilistica (Schmidt et al. 2001). Accumulation of nitrated and conformationally modified α -syn in axons has also recently been found after TBI in transgenic mice (Uryu et al. 2003).

Here, we examined whether the findings in animal TBI models of rapid axonal accumulation of proteins found in neurodegenerative diseases also occurs in human TBI. In particluar, we evaluated protein accumulation similar to that seen in AD and the synucleinopathies, including the accumulation of NF, APP, BACE, PS-1, $A\beta$, tau, and α -syn.

Materials and methods

This study was approved by the Ethics Committee of the Southern General Hospital, South Glasgow University Hospitals NHS Trust, UK.

Case Material and Preparation

Brain tissue from 18 cases following a single incident of fatal head injury was secured after full diagnostic autopsy using standardized techniques (Adams et al. 1980) by the Department of Neuropathology, Southern General Hospital, Glasgow, UK. Superficial and deep grey and white matter from the frontal lobe, temporal lobe, and brainstem was examined; however, the specific location of the tissue was unknown. None of the cases investigated had a prior history of TBI or other neurodegenerative disease. The mean +/- standard deviation age of TBI cases was 45.7 +/- 24.0 years. The survival time from TBI ranged from 4h to 5w and the post-mortem delay time was 50.2 +/- 33.6 hours.

The cause of injury was a fall in 8, a road traffic accident in 7 and assault in 3. A skull fracture was present in 14, contusions in 17 and there was an intracranial haematoma in 9. Diffuse axonal injury (Adams et al 1989) was identified in 11 (grade 3 in 4 cases; grade 2 in 2 and grade 1 in 5). Hypoxic damage was present in 15 (Graham et al. 1989) and graded as severe in 5, moderately severe in 3 and mild in 7. Brain swelling was present in 10 - unilateral in 5 and bilateral in 5, and there was internal herniation in 12 (Adams and Graham 1976). The cause of death was raised intracranial pressure in 11, pneumonia in 5, multiple injuries in 1 and systemic hypoxia in 1.

The brain of each case was collected and fixed in 10% neutral buffered formalin, then cut into slices 10mm thick and processed for paraffin embedding. Serial sections of 6 microns were cut on a Leitz rotary microtome and mounted on poly-L-lysine-coated slides for histological study.

Controls

Tissue was also secured from 6 control cases from the same institution. The mean +/- standard deviation age of control cases was 37.8 +/- 22.1 years and post-mortem delay time

was 59.2 +/- 41.3 hours. These cases had no prior history of head injury or had any evidence of structural brain damage due to pre-existing disease or injury; the cause of death in 3 was septicemia and sudden unexpected death in epilepsy (Nashef 1997; Black and Graham 2002) in the remaining 3 cases.

Immunohistochemistry

Immunohistochemistry (IHC) was performed on serial paraffin-embedded sections with investigators blinded to the cases' clinical history. Using a well-characterized panel of antibodies, we evaluated one antibody per slide for each case, resulting in the evaluation of a total of 15-20 slides per case. We chose to use several antibodies that targeted different binding sites of each protein of interest in order to provide a more comprehensive picture of the proteins present. Single-labeled IHC was carried out using the primary antibodies listed in Table 1, followed by incubation with the appropriate secondary antibody and the ABC kit (Vector Laboratories, Inc, Burlingame, CA). Visualization was achieved using DAB (Vector Laboratories, Inc, Burlingame, CA) and counterstaining with hematoxylin (Uryu et al. 2003). To evaluate the co-accumulation of proteins, we used double- or triple-labeled fluorescent immunohistochemistry (FIHC) as described elsewhere (Uryu et al. 2003; Chen et al. 2004). Briefly, tissue was incubated with a combination of primary antibodies (Table 1) followed by the appropriate fluorescent-conjugated secondary antibody. Omission of the primary antibody or application of control serum instead of the primary antibody was performed on selected sections of tissue to provide a negative control. Paraffin-embedded sections from pathologically confirmed human AD and Parkinson disease brain tissue served as positive control for tau, A β and α -syn staining.

Semiquantitative Analysis

Examination of tissue was conducted by two individuals who were both blind to the clinical circumstances of the cases. Determination of the frequency of pathological profiles containing neurofilament (NF), APP, BACE, PS1, A β , tau, or α -syn proteins was achieved by reviewing microscopic sections stained with specific antibodies recognizing the respective molecules. We followed the pathological diagnostic criteria for AD as described previously by Mirra et al, 1993 (Mirra et al. 1993). This is an observational study where sections were examined at 100x magnification and profiles were ranked semi-quantitatively as no occurrence (-), low occurrence (+), moderate occurrence (++), or frequent occurrence (+++) in a 100 μ m² field. The staining results for each protein of interest were in complete agreement, regardless of antibody recognition site; either all were positive for the protein or all were negative. Moreover, the number of profiles observed for each antibody was consistent across the tissue sections observed. Therefore, the findings presented in Table 2 are based on the consideration of all slides stained for a particular protein of interest.

Results

Recognition of axonal injury

Axonal pathology, characterized by axonal bulbs and/or swellings, was identified using antibodies to NF protein and APP.

Axonal bulbs had the appearance of discrete spherical profiles surrounded by a halo and were morphologically distinct from the varicose swellings of the axon. The bulbs ranged in size from 5-100 μ m. Only one subject in the uninjured group exhibited positive staining for NF and APP, while 15 of the 18 injured cases had extensive immunoreactivity to APP in the white matter. However, only about one-third of these cases showed positive staining for NF, an additional marker of axonal pathology (Table 2.) (Fig. 1a-d).

Accumulation and Co-localization of APP, AB, BACE and PS1

In the control group, only one subject showed APP accumulation; however, no additional protein deposits were noted. As mentioned above, almost all cases subject to TBI displayed evidence of APP accumulation within axons. In addition, BACE and PS1 staining was observed within axonal bulbs in a majority of these cases. With the exception of 2 cases, the number of profiles appeared to increase in those cases survived between 8d and 4w (Fig. 1e,f). $A\beta_{X-42/43}$ immunoreactivity within axons was noted in 13 of the total 18 injured cases (Fig 1g). $A\beta$ IHC revealed axonal bulbs positive for $A\beta_{X-42}$ but negative for $A\beta_{X-40}$. Interestingly, brain tissue from four young cases (Cases K, L, N, O) indicated rather strong axonal A β staining in their brainstem (Fig. 1h). When detected, A β plaques were limited in number and restricted to cases aged 70 years or more (Fig. 1i).

Using double- or triple-label fluorescence IHC, co-localization of $A\beta$ and its precursor protein, APP, as well as co-factors of APP processing, BACE and PS1, was examined. APP was observed to co-localize with $A\beta$ (Fig. 2a-c) or BACE and PS1 (Fig. 2d-g) within axonal bulbs following injury. $A\beta$ was also noted to co-localize with BACE and PS1 in axonal bulbs (Fig. 2h-k).

Tau and α-syn

None of the tissue from uninjured control cases stained positive for tau or α -syn proteins; and few of the injured cases stained positive for tau. However, a majority of the tissue from injured cases showed positive staining for α -syn. PHF-1 positive phosphorylated tau was noted in a small number of swollen axons and clusters of neuronal cell bodies in the cerebral cortex (Fig. 3a,b). In the same area, reactive astrocytes occasionally showed tau positive staining (Fig. 3c). Neuronal tau tended to more intensely stained than glial tau. α -syn protein was observed mostly in swollen, undulating axons as well as in bulbs, with little reactivity in the neuronal soma (Fig. 3d,e).

Summary of Principal Findings

Cases with a history of traumatic brain injury demonstrated axonal pathology in the majority of cases as shown by increased immunoreactivity to APP or NF. This compares to controls where just one case demonstrated minimal immunoreactivity. Immunoreactivity to BACE, PS-1 and $A\beta$ were all found to be increased in TBI cases versus controls.

Co-immunoreacticity between BACE, PS-1 and APP was found within axonal bulbs. In addition, within these bulbs, $A\beta$ was also found co-accumulating with APP, BACE and PS-1.

 α -syn immunoreactivity was found in two thirds of the cases, predominantly in association with axonal pathology. Only 2 cases were positive for tau which was found in both neurons and nearby reactive astrocytes. No controls had immunoreactivity for either α -syn or tau.

It is important to note that while protein accumulation was seen in many of the injured cases we evaluated, the number and location of samples that can be evaluated in an autopsy study is limited. As such, some cases graded as negative may have actually had pathologic changes in areas not examined.

Discussion

There is increasing evidence that the brains of TBI cases display many of the same pathologies associated with several neurodegenerative diseases. TBI can also induce the rapid accumulation of several proteins that compose similar pathologic aggregates found in neurodegenerative diseases such as AD, of which the most widely studied include NF proteins, APP, A β , and α -syn. Here, the accumulation of multiple proteins and their anabolic agents implicated in neurodegenerative diseases were found within damaged axons up to 5 weeks after TBI in humans. This evidence suggests that damaged axons provide a key source of proteins that may play a role in neurodegenerative processes.

Traumatic axonal injury is a common and important pathology resulting from TBI in humans (Adams et al. 1989; Adams et al. 1991; Smith DH 2000) and is frequently observed after motor vehicle accidents, falls, and assaults (Adams et al. 1982; Pilz 1983; Adams et al. 1989; Gennarelli 1993). In DAI, the axoskeleton can be severely damaged, resulting in impaired axonal transport, build-up of transported proteins, axonal swellings and bulbs (Povlishock and Becker 1985; Maxwell et al. 1997).

Axonal pathology in humans has been identified by the accumulation of NF in damaged axons from 6 hours onwards after head trauma in both animals and humans (Grady et al. 1993; Christman et al. 1994). In cases of death very shortly following injury, accumulation of NF is very limited, if present at all, in our experience. This is likely due to the fact that NF is slowly transported and accumulation is insufficient for detection in this early phase. In addition, we also note that in humans, NF immunoreactivity has a tendency to increase as survival time post-trauma elapses. This explains why in this sample we see axonal pathology as identified by NF accumulation in both fewer cases and specifically those with a longer survival period. As such, immunohistochemical detection of the fast transport APP has become a standard method of diagnosing axonal injury in human brain tissue, where axonal swellings can be identified within one hour of injury (Adams et al. 1980; Adams et al. 1989; Otsuka et al. 1991; Sherriff et al. 1994; Lambri et al. 2001; Gorrie et al. 2002; Reichard et al. 2003). The presence of acute axonal pathology is confirmed here. However, we note that within our heterogeneous population, there are cases who ultimately died secondary to raised intracranial pressure (RICP) and the vascular complications of internal herniation. Therefore, in many cases of fatal TBI it is possible to identify axonal pathology that is both traumatic and non-traumatic (infarction, related to hematomas and contusions) in origin. Axonal swellings are also seen in many other non-traumatic conditions and as part of the aging process. As so, it is appropriate to suggest that such pressure / vascular complications may have independently contributed to the neuropathological findings described (Geddes et al. 2000). Our findings also demonstrate that APP accumulation may be far more than a simple marker of axonal pathology in humans; it also may be the primary substrate for posttraumatic A β formation. In particular, in damaged axons, we observed APP accumulation along with its catalytic enzymes, BACE and PS1.To a lesser extent, other ADassociated proteins, tau and a-syn, were observed. Thus, extensive axonal damage may serve as a key reservoir of proteins implicated in neuropathologic processes. Lysis or release of accumulate proteins from damaged axons may lead to plaque formation or toxicity.

The present histopathologic findings in human TBI are consistent with findings in previous animal studies. Extensive co-accumulation of APP with $A\beta$ has been found in swollen axons in a swine model of diffuse axonal injury induced by rotational acceleration (Smith et al. 1999). More recently, in the same model, axonal $A\beta$ has been found to co-localize with its precursor protein, APP, along with the catalytic enzymes BACE and PS1 necessary to cleave $A\beta$ from APP (Chen et al. 2004).

A number of studies suggest that axons and their terminals may be a critical site of $A\beta$ production. Firstly, $A\beta$ plaques have been shown to develop in close relationship with axonal terminals in AD brains (Van Hoesen and Hyman 1990; Kamal et al. 2000; Schonheit et al. 2004). It has also been shown that APP is transported along axons by direct binding to kinesin in mouse sciatic nerves (Kamal et al. 2000). This study demonstrates that APP

operates as a kinesin-1 receptor and mediates the transport of its cleavage enzymes PS-1 and β -secretease. This in turn permits the intra-axonal generation of A β . However, this mechanism remains controversial and is contradicted by other studies (Lazarov et al. 2005). If such were the case, it is reasonable to assume that disruption of axonal transport may lead to abnormal A β generation intraxonally and potentially deposition. Indeed, Stokin et al, using an APP transgenic mouse model, demonstrated that both A β levels and deposition after disruption of axonal transport (Stokin et al. 2005). In addition, inter-axonal A β deposits formed in APP transgenic mouse brain could be reduced by anti-A β antibody therapy (Brendza et al. 2005). These studies suggest that axons can play an important role in APP processing and A β formation both intraxonally and in the generation of extracellular A β plaques. Thus, the extensive axonal damage found in TBI may provide a unique environment in which unusually concentrated co-accumulation of APP, BACE, and PS1 occurs, providing the tools to produce A β . In turn, this intra-axonal process may play a critical role in rapid A β plaque formation.

Accumulation of both tau and α -syn proteins were also found in the cases we examined; although tau was observed in fewer cases than α -syn. Based on the antibodies used, tau protein appeared to be abnormally phosphorylated and the α -syn protein was conformationally changed. These findings are consistent with the pathologies that are observed in AD brain lesions and other diseases characterized by α -syn accumulation (Fujiwara et al. 2002; Norris et al. 2004). Thus, it seems as though TBI may initiate similar processes leading to the pathological modification of these proteins that occurs in neurodegenerative disease. It is also interesting to note that both tau and α -syn are observed in the grey matter in AD brains. (Forman et al. 2004); yet here, we observed both proteins within the axons and axonal bulbs. Additionally, tau protein appeared in far fewer injured cases than did α -syn. If damaged axons provide a source of this protein, the pathological accumulation may occur over a more protracted time course than was observed here.

The present study illustrates the potential contribution of axonal injury to creating pathological protein accumulation in human brain within 4 hours – 5 weeks following TBI. Thus, it is possible that axonal injury associated with pathological protein accumulation may contribute to AD-related pathogenesis. A further understanding of the mechanistic aspects of the long-term pathophysiology of AD-related proteins in the injured brain may aid the development of interventions to halt possible TBI induced neurodegeneration.

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Figure 1.

Bright-field photomicrographs showing pathological protein accumulation associated with axonal pathology in humans. Neurofilament and amyloid precursor proteins (APP) were both found in axonal swellings and bulbs following TBI (**a-d**). Enzymatic agents of APP cleavage (BACE and PS1) were also present, although to a lesser extent (**e**, **f**). Antibodies specific for A β revealed axonal swellings (**g**) and bulbs (**g**, **inset**) positive for A $\beta_{x-42/43}$ throughout the white matter of subjects and within the brainstem of several young cases (**h**). A limited number of swellings and bulbs stained positive for A β_{x-40} . Infrequent amyloid deposits in subjects over 70 years of age were seen with A β_{x-40} (**i**). Scale bars = 50 µm.

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Figure 2.

Multiple immunofluorescent staining showing co-accumulation of APP, BACE, PS1, and A β in axons following TBI. Double-labelling revealed co-accumulation of APP and A β_{1-42} in multiple axon bulbs (**a-c**). Further immunohistochemical staining showed co-accumulation of BACE and PS-1 with APP (**d-g**) and A β_{1-42} (**h-k**).Scale bars = 50 μ m.



Figure 3.

Bright-field photomicrographs showing accumulation of tau and α -syn proteins in axonal swellings and bulbs. Of the 18 TBI cases examined, only 2 stained for tau. Tau staining was observed in axonal swellings (**a**) and bulbs (**b**) following injury, and, to a lesser extent, in glial cells (**c**). α -syn was also present within axonal swellings (**d**) and bulbs (**e**) in a majority of the injured subjects. Scale bars = 50 um.

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Summary of antibodies used for immunohistochemical analysis.

Protein	Antibody	Host	Recognition site	Dilution	Provider
Tau	17026	rabbit	Pan Tau	1:10k	CNDR
	PHF-1	mouse	pS396/pS404	1:1000	Davies [*]
	PHF-6	mouse	p+ T231	1:1000	CNDR
a-Syn	syn202	mouse	Pan synuclein	1:20,000	CNDR
	syn303	mouse	conformational a-syn	1:5000	CNDR
NF	NF-L	rabbit	NF-L	1:5000	CNDR
	RM044	mouse	Rod domains of NF-M	1:500	CNDR
	RM0217	mouse	P+ NF-H side arm, C-terminus	1:5	CNDR
ЧРР	LN39	mouse	APP	1:50	CNDR
	Karen	donkey	APP/N-terminal	1:800	Greenberg**
βk	BC05	mouse	Αβ 1-42/43	1:10,000	Suzuki ^{***}
	BAN27	mouse	Aβ 1-40	1:10,000	Suzuki
	13335	rabbit	Aβ 1-42	1:1000	CNDR
	NAB288	mouse	pan Aß	1:20,000	CNDR
	Amy117	mouse	Amyloid- 100kd protein present	1:20,000	CNDR
BACE	BACE	rabbit	BACE	1:500	Alpha Diagnostics
	BACE	rabbit	BACE/N-terminal	1:1000	CNDR
PS1	PS1	goat	N-terminal	1:100	Chemicon

CDNR = Center for Neurodegenerative Research (University of Pennsylvania), NF = neurofilament, APP = amyloid precursor protein, AB = amyloid beta, a-syn = alpha synuclein, BACE = beta amyloid cleaving enzyme, PS1 = presention-1.

* Antibody courtesy of Dr. Peter Davies: Albert Einstein College of Medicine, New York, USA.

** Antibody courtesy of Dr. B. Greenberg: Cephalon Inc., Frazer, Pennsylvania, USA.

*** Antibody courtesy of Dr. N. Suzuki: Takeda Pharmaceuticals North America Inc., Deerfiel, Illinois, USA. **NIH-PA Author Manuscript**

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Case	Age / sex	Survival	NF	APP	BACE	PS1	AMY 117	Α-β (1-42)	a-syn	Tau
V	56f	4h	1	+++++++++++++++++++++++++++++++++++++++	++++	+	-	-	++++++	1
В	47m	12h	1	1	ı	ı	-	-		1
ပ	51f	12h	1	+	+	ı	+	+	+	1
D	5m	13h	1	+	+	ı	I	I	+	1
H	57m	15h	1	‡	+	+	+	+	+	1
Ы	75m	16h	1	+	+	+		1	+	1
Ŀ	28m	18h	1	+	+	+	+	+	+	1
Н	60f	24h	1	+	+	+	+	+	+	1
Ι	65m	27h	1	+++++++++++++++++++++++++++++++++++++++	++++	+	+	+	‡	1
ſ	47m	36h	1	+	1	+	+	+	+	+
К	16m	ЪŢ	1		1	+++++++++++++++++++++++++++++++++++++++	+++++	++++	+	1
Г	15f	p8		++++	++++	++++	++++	++++	+++++	1
Μ	79f	p8			-	-	-	-	-	1
N	31m	27d	+	+	-	-	+	+	-	-
0	18m	4w	‡	+	+	+	++++++	+++++		-
Р	23m	4w	‡	+	+++++	+++++++++++++++++++++++++++++++++++++++	++++++	+++++		-
0	73f	4w		+++++++++++++++++++++++++++++++++++++++	+++++++++++++++++++++++++++++++++++++++	+++++++++++++++++++++++++++++++++++++++	++++	++++	+++++++++++++++++++++++++++++++++++++++	++
R	76m	ъ	+	+	-	-	+	+	-	-
S	15m	Control	+	+++++++++++++++++++++++++++++++++++++++	-	-	-	-	-	-
Т	26m	Control			-	-	-	-	-	-
U	54m	Control			-	-	-	-	-	-
Λ	21f	Control			-	-	-	-	-	-
Μ	73f	Control			-	-	-	-	-	-
X	38f	Control			ı		•			1

Summary of immunohistochemistry findings following TBI in humans.

Cases S-X are control subjects. (-): no occurrence of profiles in a microscopic field (~100µm2); (+): low occurrence of profiles in a microscopic field; (++): moderate occurrence of profiles in a microscopic field; (+++): frequent occurrence of profiles in a microscopic field. NF = neurofilament, APP = amyloid precursor protein, BACE = beta amyloid cleaving enzyme, PS1 = presentiin-1, AMY117 = amyloid 100kd protein, $A\beta$ = amyloid beta, α -syn = alpha synuclein.