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Targeting neurodegeneration to prevent post-traumatic epilepsy

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Abstract

In the quest for developing new therapeutic targets for post-traumatic epilepsies (PTE), identifying mechanisms relevant to development and progression of disease is critical. A growing body of literature suggests involvement of neurodegenerative mechanisms in the pathophysiology of acquired epilepsies, including following traumatic brain injury (TBI). In this review, we discuss the potential of some of these mechanisms to be targets for the development of a therapy against PTE.

Keywords

Neurodegeneration; Tauopathies; β amyloid; Neuroinflammation; Traumatic brain injury; Post-traumatic epilepsy

1. Introduction

Epilepsy is a common chronic group of neurological disorders that is characterized by the occurrence of recurrent unprovoked seizures. Post-traumatic epilepsy (PTE) accounts for up to 20% of epilepsies due to an identifiable acquired cause, i.e. “acquired epilepsies”, and follows a traumatic brain injury that initiates a process of epileptogenesis, often lasting months to years, before the first epileptic seizures are observed. The mechanisms underlying epileptogenesis are not clearly understood, but probably involve many structural and molecular alterations that alter neuronal excitation-inhibition balance and render the brain prone to generate recurrent spontaneous seizures (Avoli et al., 2016). The long duration of the pathogenesis of PTE makes it challenging to separate causal mechanisms contributing to epilepsy development from other pathological events associated with the neurotrauma which are unrelated to epileptogenesis.

Neurodegeneration is one of the characteristic brain pathologies associated with acquired epilepsies such as PTE, although the direct relationship with the development of

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spontaneous recurrent seizures is ambiguous (eg: (Brandt et al., 2006; Pitkanen et al., 2005). Nevertheless, neurodegeneration within epileptogenic regions may trigger neuroinflammation (or vice versa), network re-organization and/or cascades of molecular changes, some or all of which may contribute to the transformation from a normal to an epileptic brain network (Acharya et al., 2008). The purpose of this review is to summarise and critically analyse neurodegenerative processes as potential casual mediators of acquired epilepsy following a traumatic brain injury.

In the quest to develop antiepileptic treatments that are disease-modifying - as opposed to those providing only symptomatic relief - it is necessary to identify new biological targets that are critical to the development and maintenance of the epileptic state. In this regard, new insights can be obtained from the overlapping mechanisms of neurodegeneration that may be involved in epilepsy and neurodegenerative diseases such as Alzheimer's (AD). Patients with neurodegenerative conditions such as AD have a significantly increased risk to develop epilepsy compared to age matched controls (Abou-Khalil, 2010; Friedman et al., 2012; Nicastro et al., 2016). The accuracy of prevalence rates of epilepsy in AD are limited due to methodological difficulties of performing such studies (Friedman et al., 2012; Giorgi et al., 2017). Studies have reported that up to 64% of AD patients will undergo at least one unprovoked seizures with prevalence of epilepsy in AD patients to be up to 10 fold higher than in age matched controls (Forsgren et al., 1996; Hauser et al., 1986; Risse et al., 1990). The pathological mechanisms frequently described in neurodegenerative conditions such as hyperphosphorylated tau, β -amyloid pathology, chronic neuroinflammation are all associated with epileptic disorders. This evidence comes from studies of specimens from resected epileptogenic brain tissue from epilepsy patients who have undergone surgery for drug resistant temporal lobe epilepsy, as well as consistent robust evidences from the animal models of acquired epilepsy (Liu et al., 2016; Thom et al., 2011) The possibility of involvement of common underlying neurodegenerative mechanisms is further supported by the alteration of epilepsy outcomes on modification of these pathways (discussed in next section). However, it remains to be seen whether such alterations in epilepsy outcomes may be directly relevant to neurodegeneration mechanisms or secondary effects of modifying them. In this review, we will discuss some of the neurodegenerative mechanisms including tauopathies, amyloid pathology and neuroinflammation pathways that are relevant to PTE, mechanisms by which they may contribute to epileptogenesis or seizure susceptibility and whether modifying such mechanisms can be a target for anti-epilepto-genesis or disease-modifying treatments (Table 1). These pathways may not be the only neurodegenerative mechanisms initiated after TBI, however the ones discussed in this review have the most evidence for their relevance to epileptogenesis.

2. Tauopathies

2.1. Evidences of involvement in epilepsy

Tau is a microtubule-associated protein (MAPT) that is involved in maintaining neuronal health by affecting physiological functions including axonal transport and microtubule stabilization. For normal functioning of these processes a balance between the binding and nonbinding states of tau to the microtubules is critical. This balance is in turn regulated by

partially phosphorylated tau and controlled by kinases and phosphatase enzymes (Lee and Leugers, 2012; Morris et al., 2011). However, in a pathological environment, the balance between the non-phosphorylated and phosphorylated tau is disturbed. This leads to deposits of hyper phosphorylated tau and unbinding of tau to microtubules inducing cytoskeletal destabilization and axonal as well as synaptic dysfunctions (Lee and Leugers, 2012). These events likely contribute to the cessation of cell viability and progressing to apoptosis (Kondo et al., 2015). Apart from being associated with neurodegenerative conditions such as Alzheimer's disease and frontotemporal dementias (Rostgaard et al., 2015), an increase in hyperphosphorylated tau has also been reported following TBI (Blennow et al., 2012; Kulbe and Hall, 2017; Shultz et al., 2015). As for the evidence of involvement in the pathogenesis of epilepsy, hyperphosphorylated tau has been widely reported in clinical brain samples following surgical resection in temporal lobe epilepsy patients (Liu et al., 2017b; Puvenna et al., 2016; Sen et al., 2007; Thom et al., 2011). Similarly, pathological evaluations of brain samples from experimental models of acquired epilepsies have reported tauopathies associated with neurodegeneration and axonal dysfunction. Such evidences have been consistent across models including administration of excitotoxicants that experimentally induce status epilepticus in animals, which then leads to development of epileptic seizures or in kindling-induced seizure models (Liang et al., 2009; Liu et al., 2016; Tian et al., 2010). Likewise, alterations in tau proteins in the cerebrospinal fluid have also been reported in patients with epilepsy. Shahim et al. (2014) reported a reduction in total and phosphorylated tau within the cerebrospinal fluid in epilepsy patients following seizures. On the contrary, another study that included only patients with tonic clonic/secondary generalized seizures reported an increase in phospho and total tau in cerebrospinal fluid of the patients 48 h after seizures (Palmio et al., 2009). The discrepancy in findings could be related to the fact that the study by Shahim et al. (2014) pooled data from patients with different seizure severities including convulsive and non-convulsive seizures to non-convulsive status epilepticus. Similarly, increased phospho and total tau levels were reported following status epilepticus and was indicative of poor prognosis and a higher risk of developing epilepsy (Monti et al., 2015). Overall, these findings clearly suggest that tauopathies are related to epileptic seizures and can be observed not only in pathological evaluation of brains but could also be seen in cerebrospinal fluid.

Direct evidences come from genetic modifications in mice models that displayed alterations in the susceptibility to epilepsy development or acute seizures. Triple transgenic mouse model of AD with mutations in human amyloid precursor (hAPP), presenilin-1 and with hyperphosphorylated tau displays increased susceptibility to pilocarpine-induced acute seizures (Yan et al., 2012a). Similarly, Tg4510 tau transgenic mouse shows increased afterdischarge durations and accelerated kindling during amygdala kindling epileptogenesis (Liu et al., 2017a).

Likewise, reduction of total endogenous tau in a model of AD that expressed hAPP, reduced behavioural deficits in learning and memory as well as protected them against excitotoxicity (Roberson et al., 2007). hAPP mice with tau knockout underwent a reduced mortality, displayed less severe seizures and a longer latency to seizures following a pentylenetetrazole (PTZ) challenge in young (Roberson et al., 2007) as well as old mice (Li et al., 2014). Similar effects of reducing tau were reported in mice after kainic acid induced seizures

(Devos et al., 2013). The seizure protective effects of tau inhibition were independent of amyloid pathology and were not only relevant to chemoconvulsant-induced seizures. Protective effects of genetically inhibiting tau were also observed in genetic models of epilepsy. Genetic inhibition of Tau in *Kcna1*^(-/-) mice that displays spontaneous seizures due to lack of Kv1.1 delayed rectifier currents showed protective effects against seizure frequency and related comorbidities (Holth et al., 2013). Reduction in seizure sensitivity was also reported in the same study using two bang sensitive drosophila mutants following genetic tau inhibition (Holth et al., 2013). Moreover, tau inhibition in genetic mouse model of Dravet syndrome with mutation in voltage gated sodium channel gene prevented high mortality, and reduced the frequency of spontaneous and febrile seizures (Gheyara et al., 2014).

Some of the effects on seizure susceptibility could be the result of phenotypic changes from genetic manipulations and developmental compensatory mechanisms as a result of tau knockout. More direct evidences come from pharmacological manipulations to inhibit hyperphosphorylated tau that has shown protective effects against seizure induction as well as shown anti-epileptogenesis effects (Jones et al., 2012; Liu et al., 2016). These evidences point to clear involvement of hyperphosphorylated tau to pathophysiological mechanisms that are relevant to pathology of both genetic and acquired forms of epilepsy.

2.2. Mechanisms relevant to epileptogenesis and possible targeting for prevention of PTE

Hyperphosphorylated tau could contribute to a cascade of events such as neuronal network re-organization including mossy fiber sprouting (Decker et al., 2016; Ghosal and Pimplikar, 2011; Huang et al., 2013; Tian et al., 2010), aberrant neuronal cell migration with hippocampal granule cell layer dispersion (Kandratavicius et al., 2013), NR2B receptor mediated glutamate release (Decker et al., 2016) and neuroplasticity changes (Sotiropoulos et al., 2017) that are highly relevant to epilepsy development. However, studies showing a causal relation of hyperphosphorylated tau to above mentioned mechanisms are lacking and some or most of these effects can be considered to be associative at best. Furthermore, such neurodegenerative mechanisms could induce an inflammatory reaction that may in turn lead to enhanced excitation (see next section for details) and lead to a vicious cycle of even more cell death. An unstable/unbound microtubule assembly at the axon initial segment as an outcome of hyperphosphorylated tau is suggested to enhance excitability (Carletti et al., 2016; Matsumoto and Sakai, 1979). Though a large proportion of literature has suggested an enhanced excitability in genetic mouse models of hyperphosphorylated tau, a recent studies using either P301L tau transgenic mouse model or a triple-transgenic mouse found that there is impaired excitability in CA1 hippocampal neurons. Specifically higher threshold for action potential initiation and firing frequency as well as reduced seizure-like activity after 4-aminopyridine challenge respectively (Hatch et al., 2017; Mondragon Rodriguez et al., 2018). Such a reduced excitability in these mouse models may contribute to hippocampal and cognitive dysfunction, specifically in the case of study by Mondragon Rodriguez et al. (2018), the authors used young (1-month-old) mice and suggest that increased phosphorylation of tau in early AD disease onset may be a compensatory mechanism for increased network excitability. Overall pro-epileptogenic effects of hyperphosphorylated tau may not be only mediated by alterations in neuronal excitability but may be driven by

complex interaction of alterations in network excitability with other mechanisms relevant to epileptogenesis as described earlier.

Contribution of tauopathies to pro-epileptogenic mechanisms has provided an attractive mechanism to target anti-epileptogenesis. Tau inhibition by genetic mechanisms as discussed earlier has shown protective effects against seizures following administration of chemoconvulsants as well as in genetic epilepsy models (Gheyara et al., 2014; Holth et al., 2013; Roberson et al., 2007). However, some of the differences in findings regarding bidirectional changes in excitability in genetic tau knock out mice could be an outcome of developmental compensation or phenotypic changes in mouse model. Nevertheless, studies aiming for pharmacological targeting of tauopathies have shown clear results in terms of protecting against seizures and epileptogenesis.

Pharmacological interventions have focused both on inhibiting the phosphorylation and enhancing the de-phosphorylation pathways of tau. For a detailed discussion of pathways relevant to tau phosphorylation and dephosphorylation, the readers are suggested to refer to the book chapter by Zheng et al. (2017). Tau phosphorylation is mainly governed by the protein kinases such as glycogen synthase kinase 3 (GSK3) and cyclin dependent kinase 5 (CDK5) that induce phosphorylation in serine and threonine residues of the tau protein. Studies have suggested pharmacological inhibition of GSK3 using lithium can lead to reduced tau phosphorylation and formation of neurofibrillary tangles in mouse models with GSK3 and tau overexpression (Engel et al., 2006). Another study using old Tgtau30 mice showed reduced neurofibrillary tangles after chronic administration of lithium, though, without affecting the motor and working memory deficits (Leroy et al., 2010). Importantly, such inhibition of GSK3 pathway not only rescued tau hyperphosphorylation and neurofibrillary tangles but also inhibited neurodegeneration in AD models (Noble et al., 2005). Direct evidence of inhibiting GSK3 pathway to inhibit axonal damage from epilepsy models also exists. Liu et al. (2017b) reported a reduction in the axonal damage following inhibition of GSK3 or CDK5 by lithium and roscovitine in PTZ kindled rats (Liu et al., 2017b). Similarly, inhibition of GSK3p by 4-benzyl-2-methyl-1,2,4-thiadiazolidine-3,5-dione (TDZD-8) showed reduced neurodegeneration induced by kainic acid administration without affecting the severity of acute seizures (Bhowmik et al., 2015). Effects of targeting GSK3 pathway have only been described in AD models or post-acute seizures, though consistent findings of neuroprotection in seizure models supports potential for future studies for anti-epileptogenesis trials.

The other pathway targeted against tauopathies is dephosphorylating mechanisms of tau. In physiological conditions the phosphorylating effects of GSK3 and CDK5 are countered by tau phosphatases. Protein phosphatase 2A (PP2A) is a major tau phosphatase and its reduction in epilepsy models has been shown to be associated with hyperphosphorylated tau (Liu et al., 2016).

Sodium selenate, a potent PR55 activator (heterotrimer of PP2A), to reduce phosphorylated tau showed reduced network hyperexcitability and susceptibility to seizure induction in three different seizure models, namely— 6-Hz corneal electrical stimulation, PTZ and amygdala kindling (Jones et al., 2012). Similar seizure suppressant effects of sodium selenate were

also observed in a genetic model of Lafora disease in addition to improving the motor and memory deficits in this model (Sanchez-Elexpuru et al., 2017). While considering effects on epileptogenesis in epilepsy models, we have recently shown that inhibition of tau phosphorylation by sodium selenate inhibits epileptogenesis in three different models of acquired epilepsies (Liu et al., 2016). In this study, sodium selenate significantly delayed the progression of amygdala kindling epileptogenesis as well as reduced the number of spontaneous seizures per day and average duration of each seizure in the chronic epileptic period following kainic acid-induced status epilepticus, an effect that even persisted beyond treatment period. Similar disease-modifying effects were also observed in a PTE model following severe traumatic brain injury in rats (Liu et al., 2016). In this study, the treatment group displayed significantly fewer seizures per day, as well as a reduced average duration of each seizures. Similar to the kainic acid model, the effects on seizure frequency were sustained even after the drug treatment had ceased, suggestive of a disease modifying effect of sodium selenate treatment (Liu et al., 2016). Furthermore, targeting phosphorylated tau using sodium selenate also protects neuronal damage, brain atrophy as well as behavioural outcomes in this model that are often associated with epileptogenesis (Jones and O'Brien, 2013). Similar to genetic knock down of tau, pharmacological inhibition of tau has also been reported to improve disease pathology. Using antisense oligonucleotides to selectively decrease endogenous tau expression, DeVos et al. (2013) showed reduced seizure susceptibility in two acute seizure models with high correlation between seizure severities to tau expression. This finding supports for potentially investigating strategies to decrease endogenous tau in anti-epileptogenesis mechanisms.

Hyperphosphorylation of tau has been widely reported in experimental TBI models (Hawkins et al., 2013; Shultz et al., 2015), and also in brain tissues from human patients acutely after TBI (Albayram et al., 2017; Seo et al., 2017; Shultz et al., 2015; Thom et al., 2011). The peripheral total tau in blood of military personnel who have suffered a TBI was significantly higher than no-TBI controls, and this elevation was associated with chronic neurological symptoms (Olivera et al., 2015). Tau hyperphosphorylation has been considered a component of secondary injury following traumatic brain injury (Blennow et al., 2012; McKee et al., 2013) and is associated with decreased PP2A levels (Chen et al., 2010), the main dephosphorylating mechanism of tau. In this regard, we have previously reported a strong increase in the levels of phosphorylated tau with a concomitant reduction of PP2A activity, using a fluid percussion injury rat model of TBI (Shultz et al., 2015). Furthermore, administering the PP2A activator sodium selenate led to reduced phosphorylated tau levels, reduced brain damage including progressive atrophy of brain structures and white matter damage as well as inhibited behavioural abnormalities relevant to TBI including memory impairments, anxiety-like behavior and neuromotor performance. Reduced motor and cognitive behavioural abnormalities have also been reported after genetic inhibition of tau (Cheng et al., 2014). Importantly, inhibition of hyperphosphorylated tau following TBI has been shown to inhibit development of PTE and related brain injury (Liu et al., 2016). Despite minimal direct evidences in models of PTE, findings in models of TBI suggest involvement of tauopathies in pathological outcomes and disease progression following TBI that could be highly relevant to PTE development and potential targeting thereof.

Findings described above warrants for future studies targeting tauopathies in models of PTE. Similarly, a complete biological/biochemical mechanism for anti-epileptogenesis or neuroprotection observed by inhibiting tau-phosphorylation using sodium selenate is not established and off target effects are not clarified. In this regard, studies have delineated roles of *cis* versus *trans* isomers of phosphorylated Thr231-Pro motif in tau, suggesting that the *trans* form is physiological and the *cis* form contributes to the pathological events of hyperphosphorylated tau (Liou et al., 2003; Nakamura et al., 2012). Furthermore, a recent study from the same group targeted *cis* isoform of phosphorylated-tau in TBI mice by an antibody and reversed ultrastructural pathologies of axonal microtubules, defective cortical axonal long-term potentiation, and inhibited neurodegeneration (Kondo et al., 2015). Such strong findings establish a new strategy that can be utilized to target neurodegenerative conditions as well as PTE. At the same time, it also provides a platform for future investigations in understanding the molecular pathways relevant to other pharmacological modulators of tauopathies including sodium selenate, which has been shown to be effective against development of PTE.

3. APP and Beta-amyloid

3.1. Relevance to TBI and epilepsy

The amyloid precursor protein (APP) and its well-known proteolytic product, the amyloid- β peptide (A β), are recognized for playing a central role on the pathophysiology of Alzheimer's disease (AD), and the deleterious effects of its overexpression and clearance failure are well described in the literature (Masters and Selkoe, 2012; Mucke and Selkoe, 2012). However, the fact that APP is one of the first proteins to be overexpressed after a brain injury is a less known fact, and even though the consequences for this are still controversial, it might have an impact in the development of PTE (Blennow et al., 2012, Nature review, 2016, 2, 1–19).

Evidences for this overexpression come both from different animal models of TBI and human studies. An increase in APP immunoreactivity was observed in neurons and in astrocytes after a focal cerebral injury in rats by needle stabbing or weight drop (Lewen et al., 1995; Lewen et al., 1996; Otsuka et al., 1991). Within hours of a diffuse brain injury using the midline fluid percussion model in adult rats, APP expression was globally elevated in cortex and hippocampus; and also, as early as 1 h after an insult using the lateral fluid percussion model (Murakami et al., 1998; Pierce et al., 1996). Likewise, in a weight fall model of brainstem injury in adult rats, Yang et al. (2014) found an increase in APP mRNA levels 1 h post-impact, that peaked 3 h after injury and declined to baseline within 24 h. Similarly, as soon as 30 min post-impact in an ovine TBI model, APP mRNA was up-regulated (Van den Heuvel et al., 1999).

In humans, the findings are not different from the experimental studies and a fact worth noting is that APP immunohistochemistry is a well-validated and effective tool for diagnosing diffuse traumatic brain injury in the forensic practice and has been in use for more than 10 years (Hayashi et al., 2015). Because APP anterograde transport along axons to the synapse is disrupted in brain insult, it accumulates in the form of axonal bulbs and can be detected within hours after TBI, with some cases being detected as early as 35 min

(Hortobagyi et al., 2007; Ikonovic et al., 2004). A post-mortem study analyzed 152 patients after severe head injury with survival times ranging from four hours to 2.5 years, and found amyloid- β deposits in one or more cortical areas in 30% of the cases, and also increased APP immunoreactivity in neurons in the vicinity of amyloid- β deposits (Roberts et al., 1994). Interestingly, the amyloid- β levels in cerebrospinal and interstitial cerebral fluid (ISF) in patients after TBI correlates with the overall neurological status of the patient, in the sense that the levels increased as status improved, remained stable in stable patients and decreased when status worsened (Magnoni and Brody, 2010). This reduction in the amyloid- β ISF levels may be a consequence of the reduction of brain neuronal and synaptic activity due to injury, but it may also reflect an increased deposition of amyloid- β into insoluble aggregates; consequently, an increase in the ISF amyloid- β levels, correlating with patient recovery, might be an indication of the peptide clearance from the extracellular space although this has not been demonstrated yet (Magnoni and Brody, 2010).

Although APP upregulation after injury has been demonstrated in a number of human and animal studies, the interpretation of this fact is still under debate. Primarily, findings can be divided in two main bodies of evidence and interpreted in two different perspectives. First, that neuronal cells upregulate APP as an attempt to repair the damage caused by injury; and second, that this is a deleterious consequence of the injury itself and that APP worsens outcomes post-injury. Both perspectives will be discussed here.

The protective aspect of APP upregulation finds support in the non-amyloidogenic processing of APP. In this physiologically competing pathway, APP is cleaved within the amyloid- β region by α - and γ -secretases releasing the secreted fragment APP α (sAPP α) and an acidic intracellular domain (AICD) (for review see Muller et al., 2017). While the products of the amyloidogenic pathway, amyloid- β peptide and the secreted fragment APP β (sAPP β) may favour amyloid aggregation, sAPP α has been suggested to be neuroprotective and have a number of physiological properties that highlights its importance as a potential therapeutic target (for review see Mockett et al. (2017). Studies with transgenic models of AD have pointed towards the theory that APP processing is driven through the non-amyloidogenic pathway as an attempt to protect the brain against injury-induced cell death (Thornton et al., 2006; Van den Heuvel et al., 1999). In fact, in the APP/PS1 mouse an up-regulation in the expression of genes encoding proteins involved in A β clearance, and a reduction in A β plaques was observed after severe TBI was induced by controlled cortical impact (CCI) (Miszczuk et al., 2016). Similar results were found in a different transgenic model, the PDAPP mice, in which a reduction in A β deposit was reported even 8 months after injury while sham animals displayed increasing amyloid burden (Nakagawa et al., 1999). Likewise, aged PDAPP mice (22–24 months) with abundant A β plaques, when subjected to TBI exhibited a reduction in A β burden in the ipsilateral hippocampus compared to the contralateral site, 16 weeks post-injury (Nakagawa et al., 2000). Interestingly, mice lacking APP (APP-KO) are more vulnerable to TBI, exhibiting worst outcomes compared to the wild-type controls (WT) after a mild CCI, i.e. increased hippocampal cell loss and greater lesion volume, therefore supporting a potential protective role of APP in brain injury (Corrigan et al., 2012a). Moreover, an intracerebroventricular administration of recombinant sAPP α 15 min after a moderate injury could rescue the

damage and the functional deficits to the point that the APP-KO were no longer significantly different to the WT mice (Corrigan et al., 2012b).

Nevertheless, besides the increased risk of developing Alzheimer's by almost 2-fold after a brain injury, increasing evidence supports the fact that amyloid upregulation and deposition may also contribute to the establishment of PTE (Kenney et al., 2018). It is known that unprovoked seizures occur at rates at least 8–10-fold higher in AD patients than in the general population, and at even higher rates in autosomal-dominant and early onset cases (Amatniek et al., 2006; Mendez and Lim, 2003; Scarneas et al., 2009). Even though AD is a complex disorder and amyloid accumulation is only one of its pathological hallmarks, recent findings have strongly suggested that soluble A β oligomers when start to deposit, increase neuronal excitability and induce network reorganization that leads to hippocampal hyperactivation (Huijbers et al., 2015; Vossel et al., 2017). While hippocampal hyperactivation only occurs in the stage of mild cognitive impairment (the initial stage of AD), epileptiform activity and seizures can occur throughout the whole course of disease, and seizures that precede the onset of cognitive decline might reflect the epileptogenic potential of A β , which begins to accumulate more than 10 years before clinical signs of dementia (Bakker et al., 2015; Villemagne et al., 2013; Vossel et al., 2013).

The amyloid- β peptide may be found in different forms in the brain, as soluble monomers, oligomers, or protofibrils before aggregating into insoluble fibrils (Walsh et al., 1999; Walsh and Selkoe, 2007). The plaques are then formed outside the cell in a more advanced stage of accumulation and exert its neurodegenerative influence by a number of pathological mechanisms (Masters and Selkoe, 2012). However, it is becoming increasingly clear that the stages prior to the formation of the plaques are critical and that high levels of the soluble forms of amyloid- β disrupts synaptic transmission and impairs the excitation/inhibition balance leading to network hyperactivity (Busche et al., 2012; Mucke and Selkoe, 2012; Palop and Mucke, 2016). Indeed, a study with the APdE9 mouse model of AD reported high prevalence of seizures and increased neuronal excitability in brain slices of these mice that was attributable to the proto-fibrillar form of A β (Minkeviciene et al., 2009). The sustained resting membrane potential depolarization found in the transgenic brain slices could be mimicked by pre-incubating wild-type slices with proto-fibrillar but still soluble amyloid peptides, strongly supporting A β as a trigger for the overexcitability (Minkeviciene et al., 2009). Another study that analyzed 4 lines of transgenic mice expressing familial mutant or wild-type hAPP also found spontaneous seizure activity in cortical and hippocampal networks, although nonconvulsive (Palop et al., 2007). This aberrant excitatory neuronal activity induced by hAPP/A β triggers compensatory inhibitory mechanisms that leads to a remodeling of hippocampal circuits, consequently constraining the capacity for synaptic plasticity and contributing to network dysfunction (Palop et al., 2007).

3.2. Mechanisms of APP metabolites relevant to epileptogenesis and possible targeting for prevention of PTE

The exact mechanism of how A β induces neuronal hyperexcitability is still an open question. However, Ren et al. (2018) have proposed a mechanism based on the excessive dopamine release in the frontal cortex promoted by A β in AD and in the fact that dopamine

1 (D1) receptor is involved in A β -induced epileptic activity (Costa et al., 2016; Wu et al., 2007). By using whole-cell recordings in acute mouse brain slices they found that A β promotes dopamine release in the anterior cingulate cortex, excessively activating D1 receptors on fast-spiking interneurons and dramatically inhibiting GABA release from these interneurons (Ren et al., 2018). Therefore, by disrupting GABAergic inhibitory input, this sequence of events then leads to an excitatory/inhibitory imbalance and consequently hyperexcitability of excitatory pyramidal cells. Whether this model is also useful to explain the development of PTE after injury is something that still needs to be further investigated. Furthermore, what aggravates the impact of this amyloid cascade of events in brain injury is the fact that increased synaptic activity also seems to increase A β levels in brain interstitial fluid through synaptic exocytosis, then creating a self-sustained cycle in which amyloid is in the center (Born et al., 2014; Cirrito et al., 2005; Noebels, 2011).

Another important component that reinforces this feedback loop is inflammation and its role in the development of PTE is discussed in another section of this review. In regard to amyloid- β , its accumulation has been shown to induce microglia activation and release of pro-inflammatory mediators, which are also a big part of the neurodegeneration observed in AD (Aboud et al., 2013; Heneka et al., 2015). On the other hand, as reviewed by Webster et al. (2017) neuroinflammation per se is both a seizure trigger post-TBI and a common consequence of epileptic seizure activity. Therefore, if APP processing is driven through the amyloidogenic pathway after injury, A β toxic species could also promote epileptogenesis through inflammatory mechanisms.

Though not only A β contributes to epileptogenesis in this scenario, and it is important to consider that patients with APP duplications are also at greater risk for seizure development, as are Down syndrome patients with dementia (Cabrejo et al., 2006; Menendez, 2005). Consistently, AD mice models that overexpress APP also show hyper-excitation in individual neurons; careful monitoring of EEG often reveal epileptiform activity and spontaneous seizures in cortical and hippocampal networks (Busche et al., 2012; Busche et al., 2008; Minkeviciene et al., 2009; Palop et al., 2007). However, the problem with these AD transgenic models is that because the animals overexpress APP from birth and throughout cortical development, it is hard to delineate if the amyloid- β fragments alone, the APP and its metabolites or even both are responsible for the epileptiform activity. In this context, a recent study by Born et al. (2014) used a controllable APP-transgenic model to separate these factors and evaluate how each one contributes to network hyperactivity. It was found that genetically suppressing APP overexpression from birth until 6 weeks of age delayed the appearance of epileptiform activity, but they also found that regardless of age as long as the transgene was suppressed abnormal EEG discharges were also absent. In contrast, a reduction of A β using a γ -secretase inhibitor failed to reduce the frequency of spike-wave discharges, suggesting that APP and not A β is responsible for epileptiform activity in these particular mice.

The modulation of APP secretases as a therapeutic approach is still subject to further research. Even though Born et al. (2014) showed that pharmacological inhibition of γ -secretase failed to prevent EEG abnormalities, in the study of Loane et al. (2009) it decreased A β levels by 25%, reduced neurodegeneration and improved motor and cognitive

recovery after CCI induced TBI. This study also demonstrated that targeting β -secretase may also be useful, as mice lacking this APP processing enzyme displayed reduced cell loss and behavioural deficits after injury compared to the WT (Loane et al., 2009). Nevertheless, the non-amyloidogenic pathway, in particular the fragment sAPP α , might be worth exploiting and a better alternative as a therapeutic target in the brain injury scenario. sAPP α has been shown to have a number of physiological effects including neurotrophic, neuroprotective, neurogenic, gene expression and protein synthesis stimulation, long-term potentiation (LTP) and memory enhancement (reviewed by Mockett et al. (2017)). In regards to its neuroprotective properties, the regions of interest within sAPP α sequence were narrowed down to two domains (D1 and D6) related to their heparin binding sites and the ability of binding heparin sulphate proteoglycans (HSPGs) (Corrigan et al., 2011). As the D1 heparin binding site (APP96–110) has already proven to promote neurite outgrowth, Corrigan and colleagues administered the APP96–110 peptide intracerebroventricularly following CCI in the same APP-KO mice and could reproduce the results obtained with the sAPP α treatment (Corrigan et al., 2014; Small et al., 1994). Furthermore, improved cognitive outcomes and reduced axonal injury were also observed following APP96–110 administration in Sprague-Dawley rats after a diffuse TBI, suggesting that this peptide within sAPP α might be responsible for its neuroprotective properties (Corrigan et al., 2014).

Another potential disease-modifying therapeutic target for TBI that could impact the development of PTE is through inhibition of the c-Jun N-terminal kinase (JNK). Rehman et al. (2017) treated 2 different models of TBI mice with SP600125, a specific JNK inhibitor, and observed remarkable reductions in APP expression levels and amyloid- β production, inhibition of inflammatory responses, apoptotic neurodegeneration among other pathological features of TBI and AD-related pathology. Remarkably, treatment with SP600125 also seems to redirect APP processing from the amyloidogenic to the non-amyloidogenic pathway, without affecting A β clearance but suppressing its production (Rehman et al., 2017). Aberrant activation of JNK intracellular signaling cascade has been reported in AD patients and in AD mouse models suggesting that it might be involved in a number of neurodegenerative mechanisms associated to the disease, but this study was the first to report the benefits of targeting JNK to improve AD-related TBI outcomes (Mehan et al., 2011; Rehman et al., 2017; Shoji et al., 2000).

Lastly, targeting amyloid as a therapeutic approach in TBI involves investigating 3 main strategies: inhibition of amyloid production, prevention of its aggregation or promotion of its clearance (Schenk et al., 2012). These attempts to mitigate A β impact in the brain have been pursued through inhibition or modulation of β - and γ -secretases, blocking of A β aggregation and neutralization of oligomer toxicity, or A β immunization to promote plaque clearance. However even with a number of anti-A β treatments under clinical investigation at the moment, there has been no real progress towards plaque reduction or improvements in neurodegeneration processes (Hung and Fu, 2017). Although these approaches have been explored in the AD research field, the findings can promote insights and bolster the rationale for developing an amyloid therapy for TBI. More importantly, these treatments have not yet been explored in animal models of epilepsy, thus the relevance of whether preventing the generation of toxic amyloid species and amyloid aggregation is beneficial in terms of seizure development is still an open question. Besides, the studies herein mentioned targeting anti-

amyloid therapies in TBI models have not investigated seizure outcomes, only neurodegeneration features and cognitive improvements, highlighting the need for continued efforts in this area.

4. Neuroinflammation

Neuroinflammation is perhaps one of the most common pathological processes investigated and known to contribute to neurodegeneration following a brain trauma, as well as for its contribution to the development of epilepsies of both genetic and acquired forms (Amhaoul et al., 2016; Okuneva et al., 2015; Sun et al., 2017; Webster et al., 2017). Neuroinflammation following a traumatic injury includes many characteristic events such as activation of brain resident glial cells including microglia and astrocytes, transmigration of peripheral immune cells as well as upregulation and release of inflammatory chemokines and cytokines within local brain area as well as in distant regions including peripheral circulation (Aronica et al., 2017; Jassam et al., 2017; Webster et al., 2017). It is considered to be a major factor leading to secondary injury and neurodegeneration processes following the initial primary injury. These processes can contribute to neurodegeneration, enhancement of excitability, pruning of synapses, neuronal plasticity, neurogenesis, survival and expression of synaptic proteins on to newly formed neurons (Ali et al., 2018; Chugh et al., 2015; Iori et al., 2016; Le Feuvre et al., 2002). A more detailed evaluation of downstream pathways of how neuroinflammation can contribute to epileptogenesis has been reviewed previously (Webster et al., 2017). Of note is a recent study by Semple et al. (2017) that reported protective effect of an interleukin 1 β receptor antagonist with a reduced seizure susceptibility to evoked seizures in a mouse model of pediatric TBI along with a non-significant trend to reduced frequency of spontaneous seizures in this model. A detailed description of possibilities of targeting these in PTE has been discussed in detail in other section of this NBD supplement (Saleti et al., 2018).

Since inflammation is also one of the main repair mechanisms, here we discuss specifically whether modulating rather than complete suppression can be neuroprotective. Adverse outcomes relevant to chronic inflammation following a TBI include progression of pathological condition to neurodegenerative disorders such as AD and Parkinson's disease (PD) (Lozano et al., 2015; Utagawa et al., 2008). Chronic neuroinflammation with microglia activation and macrophage infiltration underlying pathology of these disorders involves clearing up of accumulation of β -amyloid plaques or neurofibrillary tangles. This process is compromised by an imbalance in the pro- and anti-inflammatory cytokine release with the presence of predominantly pro-inflammatory cytokines contributing to neurodegeneration (Condello et al., 2018; Laatsch et al., 2004; Laurent et al., 2018). This highlights the fact that neuroinflammation is a very heterogeneous process and the activation of immune cells in terms of their expression and release of pro- or anti-inflammatory cytokines, which has been termed to be representing a Th-1/ classical M1 or Th-2/ alternate M2 immune activation needs consideration. If unchecked, a dysregulated M1 immune response within the brain can become neurotoxic due to release of pro-inflammatory cytokines that are intended to clear up the damaged tissue. Such an immune activation in brain can be identified by the presence of markers such as pro-inflammatory cytokines including IL-1 β or TNF- α . On the other hand, an M2 response leads to the release of anti-inflammatory cytokines and growth factors

intended to promote growth and repair mechanisms, and can be identified by expression of markers such as arginase1, Ym1 and Fizz1 and release of anti-inflammatory cytokines including IL-10. These activation states of immune cells with their respective release of pro- or anti-inflammatory cytokines may also be relevant in neurodegeneration and repair after a TBI and progression to PTE.

Polarization in the microglia activation has also been reported (reviewed in (Xu et al., 2017a)) following traumatic brain injury and promoting M2/M1 ratio has shown neuroprotective effects and reduced axonal injury (Gao et al., 2016; Xu et al., 2017b). It is not known whether modification in this balance could be beneficial in protecting against PTE after a TBI. An interesting study using a genetic mouse model of progressive myoclonus epilepsy showed altered M1/M2 microglia activation balance towards an M1 phenotype before the initiation of epileptic seizures in these mice at one month age (Okuneva et al., 2015). Another recent study displayed a simultaneous expression of M1 and M2 markers during chronic epilepsy in post status epilepticus models of acquired epilepsies, though a predominantly M1 phenotype was clearly observable during the early periods after the induction of status epilepticus (Benson et al., 2015). This suggests a potential role of an M1/M2 microglia activation imbalance in pathogenesis of epileptic seizures which may be also relevant following TBI and related development of PTE. With regards to the possibility of therapeutic targeting such immune alterations, a recent study using mesenchymal stem cells to deliver interleukin-13 within the hippocampus in experimental model of epileptogenesis failed to induce a widespread induction of M2 response, and therefore a reduction in epileptogenesis (Ali et al., 2017a). Whereas, others using strategies of implanting mesenchymal stem cells have shown positive outcomes in epilepsy in association with stimulation of an anti-inflammatory immune response (reviewed in (Agadi and Shetty, 2015)). M2 promoting strategies as treatment target against PTE has not been evaluated yet, but it merits future investigations.

5. Discussion and clinical relevance

Neurodegenerative mechanisms have been widely associated with the pathophysiology of acquired epilepsies as well as following a traumatic brain injury, and represent promising targets for the development of anti-epileptogenic therapies to prevent PTE. Though, it has to be considered that neurodegeneration, at least in animal models, has not been consistently reported to be related to the development of epilepsy (Bertoglio et al., 2017; Pitkanen et al., 2005). Therefore, it must be considered how the neurodegeneration pathways discussed in this review may lead to the development of acquired epilepsies including PTE. For instance, phosphorylation of tau has been associated with a number of other molecular and pathological characteristics of acquired epilepsies such as mossy fiber sprouting, NMDA receptor subunit function, granule cell layer dispersion and alterations in neuronal plasticity (Decker et al., 2016; Kandratavicius et al., 2013; Sotiropoulos et al., 2017). Since these findings are at best associative in nature it is not possible to pinpoint to a particular mechanism linking neurodegeneration to epilepsy development. Furthermore, tau is a Microtubule Associated Protein (MAP) and it is interesting to consider whether tau is only such protein that could be targeted against PTE, or whether other relevant MAPs could be of therapeutic value. In this regard, MAP2 levels in cerebrospinal fluid have been recently

reported as a biomarker for diffuse brain injury severity and a pointer to early mortality in TBI patients (Papa et al., 2018). MAP2 is also a MAP component, localized on neuronal cell bodies and dendrites regulating spine morphology and neuronal plasticity, is known to be dysregulated both following TBI as well as in epilepsy models (Folkerts et al., 1998; Kandratavicius et al., 2013; Posmantur et al., 1996; Savina and Shchipakina, 2011; Zeng et al., 2017). Since it shares some of the dephosphorylating mechanisms with tau proteins including PP2A (Sanchez et al., 2000), a sodium selenate-mediated increase in MAP2 dephosphorylation and subsequent increase in neuronal survival cannot be ruled out. Accordingly, involvement of other MAPs such as MAPI and Echinoderm microtubule-associated protein-like protein 5 (EML5) in epilepsy pathogenesis has been suggested (An et al., 2003; Sun et al., 2015), however, a detailed discussion of these is not in the scope of this review. Similar to tauopathies, A β accumulation in the brain is known to be associated with epileptogenesis by altering dopaminergic neurons and in turn release of GABA from interneurons (Ren et al., 2018), neuronal circuit reorganization and axonal sprouting (Yan et al., 2012b) as well as neuroinflammation (Aboud et al., 2013). These suggest that the outcomes on the excitatory/inhibitory balance that leads to seizure generation is a complex interplay of several mechanistic pathways including contribution from neurodegeneration.

Genetic and pharmacological manipulations of neurodegenerative mechanisms in animal models have been suggested for possible targeting of anti-epileptogenesis trials in post-traumatic epileptogenesis. Though, in general there is scarcity of evidences in animal models of post-traumatic epileptogenesis due to a very long follow up required in post-TBI models for development of epilepsy and a relatively low incidence (20–30%) in adult TBI rodent models. This renders undertaking adequate, statistically powered, studies challenging in terms of expenses as well as the logistics of performing such studies. However, the role of neurodegenerative mechanisms in other acquired epilepsy models and TBI outcomes strongly suggest it is important to invest in this line of research.

In this regard, two of the pathways are particularly promising – inhibition of tau phosphorylation by sodium selenate and inhibition of pro-inflammatory interleukin 1 β . Targeting tauopathies with the use of sodium selenate has shown effectiveness in inhibiting epileptogenesis in three different models of acquired epilepsy, including the one initiated following TBI (Liu et al., 2016). Sodium selenate has already been reported to be safe during phase I and phase II trials for non-epilepsy conditions (Corcoran et al., 2010; Malpas et al., 2016), and so could rapidly be translated into human anti-epileptogenesis studies post-TBI. Similarly, effectiveness of modifying interleukin 1 β signaling has shown promising effects in models of acquired epilepsies (Fukuda et al., 2009; Noe et al., 2013) including in a model of TBI (Semple et al., 2017). Interleukin 1 β antagonist Kineret is already commercially available for treatment of rheumatoid arthritis. In addition, clinical case reports have also suggested promise of improving outcomes in epilepsy patients (Desena et al., 2018; Kenney-Jung et al., 2016). Further validation of these in animal models of PTE is being undertaken in the EpiBioS4Rx (Epilepsy bioinformatics study for antiepileptogenic therapy) Center Without Walls funded by NIH, which will test new therapies to prevent PTE, in a multi centered preclinical trial scenario (discussed in detail in another work in this special issue: Saletti et al 2018). Overall, we provided several neurodegenerative mechanisms that may be relevant to development of PTE. Some of those may still be at the

stage of proof of concept evidence whereas some others have already shown clear potential for progressing into clinical trials targeting the development of PTE.

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Table 1

Summarizing the neurodegenerative mechanisms including Tauopathies, amyloid pathology and neuroinflammation in patients and animal models with TBI.

Model	Mechanism/s investigated	Intervention/s	Outcomes	Reference
TBI patients	Hyper-phosphorylated-Tau	Not evaluated	Increased Hyperphosphorylated-Tau	(Shultz et al., 2015)
Repetative TBI in patients	Hyper-phosphorylated-Tau	Not evaluated	Increased phosphorylated tau in rTBI patients that were similar to observation in temporal lobe epilepsy patients	(Puvanna et al., 2016)
TBI patients	Hyper-phosphorylated-Tau	Not evaluated	Phospho-Tau associated with TBI history but not seizures	(Thom et al., 2011)
Traumatic encephalopathy patients	PP2A activity	Not evaluated	Decreased PP2A activity and associated genes in patients	(Seo et al., 2017)
Military personnel with TBI	Total Tau	Not evaluated	Increased total tau in blood that related to increased post concussive neurological symptoms	(Olivera et al., 2015)
TBI patients	Cis-Tau	Not evaluated	Cis-Phospho-Tau increased in cortical axons and CSF and correlated with axonal injury and clinical outcomes	(Albayram et al., 2017)
rTBI in mice		Specific neutralizing antibody against cis phospho-Tau	Attenuates neuropathology and brain dysfunction.	
Lateral fluid percussion injury model in Long-Evans rat	PK55 and PP2A levels regulating Hyper-phosphorylated-Tau	Tau phosphatase activator-sodium selenate	TBI reduced PK55 and PP2A activity and increased phosphor-Tau expression that was reversed by sodium selenate. Reduced brain damage and improved behavioural outcomes.	(Shultz et al., 2015)
Lateral fluid percussion injury model in Long-Evans rat	Hyper-phosphorylated-Tau	Tau phosphatase activator-sodium selenate	Reduced epileptic seizure frequency and duration in treated rats with effects observable even following washout period	(Liu et al., 2016)
Parasagittal FPI model in rats	Hyper-phosphorylated-Tau	Not evaluated	Increased oligomeric and Phospho-Tau	(Hawkins et al., 2013)
Mechanical compression in rats	Hyper-phosphorylated-Tau and microtubule associated protein-2	Not evaluated	Increased phosphorylation of Tau and MAP2 with decrease PP2A activity within minutes	(Chen et al., 2010)
Mild rTBI in mice	Total Tau	Complete ablation/partial reduction of Tau	Reduced spatial learning and memory deficits along with reduced level of axonopathy	(Cheng et al., 2014)
Surgical excision of human temporal cortex after TBI	APP overexpression	Not evaluated	Increase in APP levels after TBI in patients.	(Ikonomovic et al., 2004) (Roberts et al., 1994)
Post-mortem study with 152 patients that had severe TBI				
Needle stabbing or weight drop in rats	APP overexpression	Not evaluated	Increase in APP levels after TBI in different animal models.	(Lewen et al., 1995)/ (Otsuka et al., 1991) (Murakami et al., 1998) (Yang et al., 2014)
Midline fluid percussion model in rats				

Model	Mechanism/s investigated	Intervention/s	Outcomes	Reference
-Weight fall model of brainstem injury in rats				(Van den Heuvel et al., 1999)
Ovine TBI model				
Severe controlled cortical impact in APP/PS1 mice	APP processing after injury	Not evaluated	Up-regulation in the expression of genes encoding proteins related to A β clearance.	(Miszczuk et al., 2016)
Controlled cortical impact in young PDAPP mice	APP processing after injury	Not evaluated	Reduction in A β deposit after injury while sham animals displayed increasing amyloid burden.	(Nakagawa et al., 1999) (Nakagawa et al., 2000)
Controlled cortical impact in aged PDAPP mice			Reduction in A β burden in the ipsilateral hippocampus after injury in mice with abundant plaques.	
Mild controlled cortical impact in APP-KO mice	APP role after injury	Not evaluated	APP-KO mice exhibit worst outcomes after injury compared to wild-type controls.	(Corrigan et al., 2012a)
Moderate controlled cortical impact in APP-KO mice	APP role after injury	Intracerebroventricular injection of sAPP α . 15 min after injury	sAPP α treatment rescued damage and functional deficits in APP-KO mice.	(Corrigan et al., 2012b)
Controlled cortical impact in mice	APP metabolites	γ -secretase inhibitor	Reduced A β levels by 25%, reduced neurodegeneration and improved motor and cognitive recovery after TBI.	(Loane et al., 2009)
Controlled cortical impact in APP-KO mice	APP metabolites	Intracerebroventricular injection of APP 96-110 peptide	Improvement of outcomes post-injury similar to the treatment with sAPP α .	(Corrigan et al., 2011)
Diffuse TBI in Sprague-Dawley rats	APP metabolites	Intracerebroventricular injection of APP 96-110 peptide	Reduced axonal injury and improved cognitive outcomes.	(Corrigan et al., 2014)
Feeney's weight drop model and repetitive mild TBI in mice	APP metabolites	SP600125 (JNK inhibitor)	Reduction in APP expression levels and A β production, inhibition of inflammatory responses and apoptotic neurodegeneration.	(Rehman et al., 2017)
Pediatric TBI model in mice	Interleukin-1 β	Kineret (Interleukin-1 receptor antagonist)	IL-1Ra treated mice displayed improved spatial learning behavior, less cortical damage and showed fewer evoked seizures compared to controls	(Semple et al., 2017)
Severe controlled cortical impact in mice	M1/M2 polarization Neuroinflammation	Intracerebral injection of human neural stem cells	No change in lesion size but reduced accumulation of amyloid precursor protein, transition of microglia towards M2 phenotype	(Gao et al., 2016)
Controlled cortical impact in mice	M1/M2 polarization Neuroinflammation		Improved M2/M1 ratio with increased anti-inflammatory cytokine expression, reduced total microglia/macrophage expression, reduced peripheral invasion of immune cells, reduced apoptosis and improved behavioural deficits	(Xu et al., 2017b)