Role of metal ions in the cognitive decline of Down syndrome

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INTRODUCTION

This review will discuss the role of intrinsic metals in cognitive decline in people with Down syndrome (DS). DS, caused by trisomy of whole or part of human chromosome 21 is the most common mental impairment with an incidence of about 1 in 700 live births (Epstein, 1995). Characteristics of DS individuals have been described such as stunted growth, mental impairment, congenital disorders of multi-organ systems such as the heart, haematological, musculoskeletal, thyroid, gastrointestinal, dental, nervous and immune systems, and having a higher incidence of diabetes and age-associated diabetes (Burch and Milunsky, 1969; Peiris et al., 2012). Life expectancy of people with DS has increased from 12 years in the 1940s to 60 years at the present time, largely due to enhanced medical and social care (Contestabile et al., 2010). Because of the increased longevity, it has become evident that by the age of 40 (Mann et al., 1990), all individuals with DS have the cognitive decline and the neuropathology seen in Alzheimer's disease (AD).

AD is the most common form of dementia. It was first described by a German psychiatrist and neuropathologist named Alois Alzheimer in 1906 (Alzheimer et al., 1995). One of the earlier symptoms is short term memory loss. As the disease progresses long-term memory loss, confusion and mood swing occur. Sporadic AD and DS share similar neuropathological features (Mann, 1988), these include the enlargement of early endo-somes in certain neurons, deposition of A β plaques, the presence of tau-containing neurofibrillary tangles and the degeneration of basal forebrain cholinergic neuron (BFCN). The pattern of

Down syndrome (DS), caused by trisomy of whole or part of chromosome 21 is the most common mental impairment. All people with DS suffer from cognitive decline and develop Alzheimer's disease (AD) by the age of 40. The appearance of enlarged early endosomes, followed by Amyloid βpeptide deposition, the appearance of tau-containing neurofibrillary tangles and basal forebrain cholinergic neuron (BFCN) degeneration are the neuropathological characteristics of this disease. In this review we will examine the role of metal ion dyshomeostasis and the genes which may be involved in these processes, and relate these back to the manifestation of age-dependent cognitive decline in DS.

Keywords: Down syndrome, Alzheimer disease, iron, copper, zinc, calcium

pathology is similar but the symptoms appear earlier in individuals with DS (Ikeda et al., 1989). In DS, enlarged endosomes in neurons are seen as early as 28 weeks of gestation (Cataldo et al., 2000). The presence of enlarged endosomes precedes $A\beta$ peptide deposition which appears at around 12 years of age in the form of diffuse plaques, followed by mature AB plaques when the individuals are in their 30's (Lemere et al., 1996). In a review of the literature, it was reported that most studies have found that tau-containing neurofibrillary tangles occur in the BFCN along with neuronal degeneration, gliosis and dementia at ages of between 35 to 45 years (Wilcock and Griffin, 2013). An immunohistochemical study of the brain tissues from postmortem DS individuals showed that in earlier years of life (37–38) brain neuron loss was present and this progressed with age (over 50s) with abundant, mature amyloid deposits. Taken together, these observations suggest that in DS there is an accelerated progression occurring through the different stages of AD-like neuropathology.

There are 127 known genes, 98 predicted genes and 59 pseudo genes located on chromosome 21 (Hattori et al., 2000). It is putatively thought that having an extra copy of one or more of these genes could be responsible for the manifestation of AD-like neuropathology and cognitive decline in DS. Some genes of interest are: Amyloid precursor protein (*APP*), Down syndrome candidate region 1 (*DSCR1*), Intersectin 1(*ITSN1*), Superoxide dismutase 1(*SOD1*), Beta-site APP-cleaving enzyme 2 (βsecretase or *BACE 2*), S100 calcium binding protein $\beta(S100\beta)$.

In the next section, how these genes may be involved in cognitive decline of DS will be discussed.

GENES POTENTIALLY INVOLVED IN COGNITIVE DECLINE OF DS

AMYLOID PRECURSOR PROTEIN (APP)

Studies have suggested that over-expression of the APP is the key element for the manifestation of AD in DS (Schupf and Sergievsky, 2002). A case study where an individual had partial trisomy 21 that did not include an extra copy of the *APP* gene confirmed this concept (Prasher et al., 1998). This individual showed some characteristics of DS but did not suffer from AD and the associated cognitive decline.

Amyloid peptides (A β 1–40/42) are cleavage products of the APP. These form β -sheet rich aggregates and soluble oligomers. The cleavage products are generated by the proteolytic enzymes β secretase and γ secretase. There is also an alternative pathway where α -secretase cleaves APP. The products of this pathway do not include A β peptides and aggregates do not form (Vassar et al., 1999). The gene encoding β secretase 2 is located on chromosome 21, therefore, an extra copy would potentially contribute to the production of higher levels of toxic A β , hence formation of amyloid plaques early in DS.

However, this extra dosage of APP and BACE may not be required to produce the AD-like phenotype in DS. This idea can be derived from observations from an individual who had an over-expression of APP, but exhibited normal levels of BACE (Cheon et al., 2008). In a recent study, polymorphisms in BACE2 were shown to be a major factor for the age of onset of AD in the DS population (Mok et al., 2014) suggesting a significant role for BACE2 in the development of AD in DS.

In addition, it was shown that the amyloid truncated A β peptides (A β 9–42 and A β 17–42) form ion channels by formation of a β -barrel secondary structure (β strand-turn- β strand) that alters calcium (Ca²⁺) homeostasis providing additional avenues for the manifestation of AD (Jang et al., 2010). An alternative mechanism may be that N-terminal truncated amyloids are neurotoxic because they form fibrils that aggregate more readily (Pike et al., 1995). These studies suggest that it is not only A β but smaller truncated amyloid peptides which are also involved in the development of AD.

AD research has focused heavily on APP. However studies have suggested that APP may not be the only factor involved in the manifestation of AD. This was demonstrated in mice where the APP protein was over-expressed in a mouse model of DS (Ts65Dn). No change in the endosomal phenotype was observed indicating other factors are necessary for this alteration (Cataldo et al., 2003). As mentioned above, there are other genes that could contribute to the development of ADlike neuropathology that will be discussed later in this review. Although APP is necessary to develop the disease, it appears that it alone is not sufficient to cause it. This is why it is interesting to study DS as it narrows the search done to trisomy genes.

Importantly, amyloid plaques are formed as the result of inappropriate interaction between biometals (iron, copper and zinc) and beta-amyloid. Indeed, increased levels of zinc promote A β precipitation (Bush et al., 1994). There is also increased levels of iron in AD patients (Gerlach et al., 1994). These observed increases in the levels of metals in AD give rise to the questions

of why, how and when these increases occur? This so-called metal theory of AD and DS will be discussed later in this review.

DOWN SYNDROME CANDIDATE REGION 1

Studies showed that DSCR1 mRNA expression was increased in post-mortem brains of AD patients compared with agedmatched non-AD controls (Ermak et al., 2001). DSCR1 is a regulator of calcineurin and in some cases has been shown to inhibit calcineurin signaling pathways (Fuentes et al., 2000). Over-expression of DSCR1 contributes to the formation of neurofibrillary tangles and A β deposition (Ermak et al., 2011). As discussed earlier, formation of neurofibrillary tangles is the first step of memory loss. Hence, having an extra copy of DSCR1 in DS could contribute to earlier formation of neurofibrillary tangles and premature cognitive decline. The formation of tangles could be facilitated by the enhanced ability of over-activate calcineurin to dephosphorylate tau but the mechanism has not been fully elucidated (Ermak et al., 2001; Ma et al., 2004).

INTERSECTIN 1

Another potential gene of interest on chromosome 21 is *ITSN1*. This gene appears to be involved in metal homeostasis since it is involved in receptor mediated endocytosis and one of its cargos is the transferrin/transferrin receptor complex required for iron transport into the cell. Transferrin (and its iron cargo) is transferred from the blood into cells by the formation of a vesicle (clathrin-coated pit) that internalizes the complex. First the recruitment of adaptor protein complex 2 (AP-2) to the plasma membrane occurs, and then clathrin is recruited to the cytoplasm where AP-2 has been recruited. The plasma membrane forms a vesicle by budding inward. The receptors and bound ligand (transferrin) are thus taken up by the cell. ITSN1 is involved in the recruitment of AP-2 and the formation of clathrin-coated pits. Having an extra copy of the *ITSN1* gene in DS could alter receptor trafficking and consequently iron transport.

METALS

THE METAL THEORY

As discussed above, excess APP and/or its cleavage products is necessary for the development of AD but, it is not sufficient to cause it. In addition, clinical trials targeting A β (by anti-A β antibodies) have failed to treat AD (Greenberg et al., 2003). Bush et al. (on the basis of the studies done by them and others) have proposed that a number of proteins involved in neurodegeneration (A β , A β PP, tau, presenilin, and β -secretase1) fail in their ability to regulate metals in the AD brains (Finefrock et al., 2003; Bonda et al., 2011; Bush, 2013), and consequently these proteins are overwhelmed by the increased levels of these metals. This leads to accumulation of extracellular zinc and copper in amyloid, and an accumulation of intracellular iron in neurons (Bush, 2013).

Iron

Iron is essential for normal neurological function. It is required for the synthesis of myelin and neurotransmitters (Piñero and Connor, 2000; Bush, 2013). It is known that free iron accumulates in the brains of AD patients, and that iron transport and storage are disrupted (Sayre et al., 2000). Another study showed that iron stimulates the production of hydroxyl radicals from hydrogen peroxide, and hence exacerbates oxidative stress. Iron was increased in all areas of the AD brains compared with controls and transferrin was increased in the frontal cortex of AD brains. The transferrin/iron ratio which is indicative of iron mobilization was reduced in AD brains, indicating a disturbance of iron metabolism in the AD population (Loeffler et al., 1995).

Iron transport into the brain is accomplished by Transferrin (Tf). Tf is a glycoprotein that carries iron in the blood and regulates iron homeostasis. As mentioned above, ITSN1 is required for iron internalization. It has been reported that there is an increase of Tf in brains of DS individuals compared with controls (Leveugle et al., 1994). An extra copy of *ITSN1* could be the reason for this elevated level of Tf in DS brains, and this level increases with age in DS. The increased level of Tf is also evident in the amniotic fluid obtained from women carrying DS foetuses (Perluigi et al., 2011), suggesting iron-mediated damage starts during pregnancy.

Ferritin, the iron storage protein is available in the cytosol to bind the iron that has been endocytosed into the cell. Excess iron can be transported out of the cell by ferroportin (Fpn). For this to occur, a ferroxidase enzyme is required. In the glia, ceruloplasmin is the known ferroxidase (Klomp et al., 1996) but ceruloplasmin is not produced by neurons. Instead, APP has been implicated in the process of iron export in neuronal cells. This occurs through the stabilization of Fpn (Duce et al., 2010). Interestingly, excess dietry iron fed to APP-knock-out mice induced iron accumulation and damaged the neocortex (Duce et al., 2010).

Iron regulatory protein (IRP) is a cytoplasmic protein that regulates iron uptake, storage and usage in response to the concentration of cellular iron. The level of iron regulatory protein (IRP-2) in the whole brain was the same in both AD patients and controls. However IRP-2 is co-localized to the specific areas with AD neuropathology (Farrar et al., 1990; Smith et al., 1998), suggesting the regulation of iron uptake, storage and usage changes in specific parts of AD brains which leads to the development of the AD neuropathology. The role of IRP-2 and indeed iron in the patho-biology of AD has been well established, whereas in DS this needs to be investigated.

Tau protein stabilizes microtubules which are involved in cellular transport. In AD, excess tau phosphorylation causes formation of tau clumps (neurofibrillary tangles). This leads to a break down of nutrient transport and death of the neurons which is believed to be the first step of memory loss. The appearance of tau-containing neurofibrillary tangles is one of the later neuropathological hallmarks in AD brains. Tau can bind to iron and this could contribute to tau aggregation (Adlard and Bush, 2006). Tau protein is needed for iron-export as it was shown that iron accumulated in tau-knock-out mice (Lei et al., 2012). Therefore, excess iron levels in the brain could contribute to memory loss as the consequence of an increased level of tau aggregation.

Copper

The brain is enriched with copper. Copper has been localized in mitochondria, synaptosomes and myelin (Nalbandyan, 1983). In a normal nervous system, copper plays a role in regulating neuronal excitability (Kardos et al., 1989), myelination, iron metabolism and the function of copper containing enzymes such as superoxide dismutase (SOD), dopamine- β monooxygenase and tyrosinase (Nalbandyan, 1983). As mentioned above, the *SOD1* gene is located on chromosome 21. An extra copy of *SOD1* in DS might contribute to a dysfunctional SOD1 enzyme. SOD is a copper and zinc binding enzyme which is responsible for destroying free superoxide radicals in the body.

Studies have also shown that APP knockout mice have higher levels of copper in brain and liver (White et al., 1999) while the APP-overexpressing transgenic mice have reduced levels (Maynard et al., 2002) indicating that APP acts as a major regulator of neuronal copper homeostasis. On the other hand, a recent study showed that copper promotes APP trafficking through the secretory pathway (Acevedo et al., 2014). Collectively, these studies suggest there is a two way relationship between copper and APP.

It is known that copper binding to APP reduces the production of A β *in vitro* (Barnham et al., 2003). As DS individuals have an extra copy of the *APP* gene, it may follow that they would have reduced levels of copper in their brains and hence increased production of A β leading to the formation of amyloid plaques. This has yet to be investigated.

It has also been shown that in AD brains, there is an overall deficiency of copper (Cater et al., 2008) and an increased level of extracellular copper in amyloid (Bush, 2013) suggesting a change in the homeostasis of copper in AD. Excess cellular copper can compromise cell viability by acting as a pro-oxidant and generating toxic reactive oxygen species via Fenton-type reactions, involving copper ions and hydrogen peroxide. Therefore, cellular copper homeostasis must be tightly regulated (Huang et al., 1999) to maintain healthy brain function.

Zinc

Zinc is essential for the function of numerous enzymes and transcription factors. In the normal brain, zinc is bound to membrane-bound metalloproteins, or loosely bound within the cytoplasm to proteins and enzymes as well as being in synaptic vesicles that are enriched with zinc (exceeding 1 mmol/L in concentration) (Frederickson et al., 2000). During synaptic transmission, high concentrations of zinc are released into the synapse from mossy fibers (Sensi et al., 2011). Zinc regulates ion channels and transmitter receptors such as α -Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) and N-methyl-D-aspartate (NMDA) receptors which are implicated in synaptic plasticity and memory consolidation (Sindreu and Storm, 2011). Therefore, any change in homeostasis of zinc can have an impact on learning and memory.

Within the ectodomain 2 (E2) of APP, a metal binding site has been identified which binds competitively to zinc or copper (Dahms et al., 2012). Both zinc and copper can interact with A β to form aggregates (Adlard and Bush, 2006). An increase in level of extracellular zinc in amyloid of AD has been observed (Bush, 2013). Earlier studies demonstrated that a small increase in brain zinc concentration (>3 micromolar) increased the adhesiveness of A β (Bush et al., 1994) and changed A β metabolism. Intriguingly, the area of the brain with the highest level of zinc, the cerebral cortex (Frederickson et al., 1983), exhibits the most severe pathological lesions of AD (Hyman et al., 1986).

Zinc transporters (ZnT) are also crucial to maintain memory and cognitive function. There are 10 ZnT known to date. Zinc uptake into synaptic vesicles needs zinc-transporter-3 (Cole et al., 1999). It has been demonstrated that zinc-transporter-3 knockout mice develop amyloid pathology characteristic of AD (Adlard et al., 2010) supporting a role for the zinc transporter in the manifestation of AD. Another study showed that zinc-transporter-6, which is located in the Golgi apparatus, is involved in the accumulation of zinc (Lyubartseva et al., 2010). The role of ZnT is yet to be investigated in DS.

Calcium (Ca²⁺)

Calcium plays an important role in the central nervous system. It acts as a cofactor, second messenger and signaling molecule, and a coenzyme when part of a protein. It is also part of excitatory function of neuronal cells, being involved in voltage-gated calcium ion channels. It has been shown that voltage-gated calcium ion channel activity in a mouse model of DS (trisomy 16 mouse; Ts16) was significantly higher than the wild type (Galdzicki et al., 1998). This increased activity could potentially lead to altering calcium homeostasis in the brains of DS individuals.

Human chromosome 21 has two genes of interest that modulate calcium; the *S100* β gene stimulates calcium influx (Mattson et al., 1993a,b) and the *DSCR1* gene, which is a regulator of calcineurin and in some circumstances, can inhibit calcineurin signaling pathways. It has been shown that calcineurin regulates Ca²⁺ pumps and exchangers to maintain Ca²⁺ homeostasis (Stark, 1996). Having an extra copy of *DSCR1* could potentially change calcium homeostasis in the brains of people with DS by influencing the activity of calcineurin, leading to formation of neurofibrillary tangles and consequently memory loss.

In AD, AB binds to neurons in close proximity to the NMDA-R which triggers NMDA-R- mediated calcium influx and alters calcium homeostasis (De Felice et al., 2007). Another report suggesting altered calcium homeostasis contributed to AD came from a study on L-type voltage gated calcium channels (LTVGCC) (Anekonda et al., 2011). In this study MC65 neuroblastoma cells were exogenously transfected with APP under the control of a tetracycline-responsive promoter. Upon the withdrawal of tetracycline, production of APP and Aß fragments was observed along with up-regulation of LTVGCC, leading to increased calcium influx (Anekonda et al., 2011). Interestingly the use of calcium channel blockers prevented the neurotoxicity of AB. It has been suggested that the neurotoxicity of $A\beta$ is due to the presence of elevated calcium ions, resulting in increased responses to excitatory amino acids such as glutamate (Mattson et al., 1993a). Due to these increased responses, glutamate receptors are over activated which result in dendritic pruning, increased immunoreactivity of tau, and an accumulation of filaments. These are all characteristics of neurofibrillary tangles of AD in DS and non-DS individuals.

CONCLUSION

An extra dosage of the genes on chromosome 21 could contribute directly or indirectly (by changing metal homeostasis) to the

manifestation of AD-like neuropathology in DS. There might be more genes involved in these processes other than the ones discussed in this review. Although some studies have been conducted on cognitive decline in DS individuals, the molecular mechanisms involved in this cognitive decline remain unknown. More studies are required to investigate these mechanisms and thus provide avenues to explore for the prevention of the cognitive decline characteristic of AD in DS and in the non-DS population.

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