



Opinion Article

Is the Regulation of Microtubule Stability at the Crossroad between Aging and Disease of Dopaminergic neurons?

Cartelli Daniele*¹ and Cappelletti Graziella^{1,2}

¹Department of Biosciences, Università degli Studi di Milano, Milano, Italy

²Center of Excellence on Neurodegenerative Diseases, Università degli Studi di Milano, Milano, Italy

Abstract

Over the time, it is becoming clear that microtubule system plays a fundamental role not only during the development of central nervous system but also during brain aging and neuronal death. In particular, the fine modulation of microtubule stability and tubulin posttranslational modifications is crucial for neurons as its failure induces brain damage. Basing our idea on the facts that dopaminergic neurons possess a peculiar architecture, strictly dependant on microtubule integrity, and that most of the experimental models of Parkinson's disease undergo microtubule destabilization before dopaminergic cells loss, here we propose that the regulation of microtubule stability can be the Achilles' heel of this neuronal population. Therefore, we hypothesize that failure of the modulation of microtubule system could represent the crossroad between normal brain aging and neurodegenerative processes.

Keywords: Dopaminergic neuron; Microtubule stability; Parkinson's disease; Tubulin acetylation

Microtubules (MTs) are built up by $\alpha\beta$ tubulin dimers, which assemble into non-covalent polymers with an intrinsic resistance to bending and compression and capable to alternate slow polymerizing phases to rapid shrinking ones, a behaviour called "dynamic instability" [1]. The proper regulation of MT stability (i.e., more stable MTs display a longer half-life) is fundamental for the development, the survival and the aging of neurons, since both hyper-stable and overly dynamic MTs affect neuronal functions eventually leading to cell death [2]. α and β tubulin Posttranslational Modifications (PTMs) mark dynamic or stable MTs, which are found in specific neuronal compartments and whose stability changes during neuronal differentiation and aging [3]. Indeed, tyrosinated α tubulin is associated to highly dynamic MTs, which are abundant in young neurons and at the

presynaptic bouton of mature neurons. During neuronal maturation MT stability increases and MTs inside the axonal shaft accumulate PTMs which extend their half-life, either through a direct stabilization of the MT lattice [4] or by governing upstream events, such as the recruitment of MT-interacting protein [5] or by protecting MTs from depolymerizing kinesins and severing enzymes [6,7]. Nevertheless, the stability of MTs has to be maintained under a physiological threshold, since their excessive stabilization can eventually lead to MT bundling and block of axonal transport [8]. As many of the axonal MTs are long-lived polymers which persist for the entire life of the neuron, their aging is a critical event which potentially impacts on the neuron lifespan. Indeed, MTs are structures with an intrinsic self-repair capability [9] but, due to defects accumulated during growth [10] and to their ends which become more tapered during time [11], the older MTs are more susceptible to catastrophes than younger ones [12,13]. Interestingly, tubulin PTMs are potential regulators of this process as much as, for long time, acetylated MTs were referred almost synonymously to old and stable MTs. Indeed, among its several biological functions [14], it has been shown that acetylation accumulates during MT aging and it has been proposed that the MT-acetylase α Tubulin Acetyl Transferase1 (α TAT1) can act as a clock for MT lifetimes [15], due to its preference for MTs and its low catalytic rate. Nevertheless, other candidates may be responsible for minor tubulin acetylation *in vivo*, including Elongator Complex Protein E1p3, which acts as tubulin acetylase, controlling the development and the migration of cortical neurons [16]. Furthermore, the age-dependent decrease of the activity of sirtuin 1, a widespread deacetylase which also acts on MTs, could lead to increased acetylation of α tubulin in the cerebellum of aged mice [17]. These data suggest a functional link between MT acetylation patterns and brain aging, evidencing how the regulation of MT stability is critical during normal neuronal aging and supporting the concept that its failure may be a reliable candidate in causing neurodegenerative processes. Indeed, during the last years, an increasing body of evidence indicated the potential role of defective regulation of MT stability in neurodegenerative disorders [18] and, in particular, in Parkinson's Disease (PD) [19,20].

PD is the second most common neurodegenerative disorder caused by the death of the dopaminergic neurons in the *Substantia nigra* [21] whose anatomy, with an average arborisation of 4.6 meters in humans [22], makes them particularly dependant on axonal transport which, in turn, depends on MT integrity and whose failure has been associated to the early phases of PD [23]. Although the identity of the real culprit of the pathology is still debated, in the recent years it is becoming ever clearer that MT destabilization is an early event in experimental and human models of PD, and many of the studies reported below strongly suggest that it can be followed by the hyper-stabilization of MTs, which could lead to detrimental side effects. Indeed, during the last decade, we have demonstrated that MPP⁺, the metabolite of the PD-inducing toxin MPTP, acts as a catastrophe promoter, leading to the earlier MT destabilization and to a later excessive accumulation of tubulin PTMs associated to stable MTs (detyrosination and acetylation), *in vitro* as well as *in vivo* [24-26]. Similarly, 6-hydroxydopamine elicits significant alterations in MT dynamics,

*Corresponding author: Daniele Cartelli, Department of Biosciences, Università degli Studi di Milano, Via Celoria 26, Milano, 20133, Italy, Tel: +39 3334920793; E-mail: danielle.cartelli@gmail.com

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including reduction in MT growth rate and overload of a tubulin acetylation [27]. Furthermore, MT fragmentation has been observed in various dopaminergic neuronal models exposed to various PD-inducing neurotoxin [28-30 and Casagrande et al., personal communication], and Ren and colleagues clearly demonstrated that dopaminergic neurons are much more susceptible to MT destabilization than other neuronal types [28]. Moreover, some and perhaps the most important PD-related proteins interact with the MT system modulating its assembly and stability. We recently demonstrated that wild type α -Synuclein is a MT dynamase which regulates MT nucleation and dynamics and whose PD-associated mutations induce tubulin aggregation [31]. Both parkin and Leucine Rich Repeat Kinase 2 (LRRK2) interact with and stabilize MTs [32,33] and, notably, induced Pluripotent Stem Cells (iPSC)-derived neurons obtained from patients carrying mutations in parkin and LRRK2 display MT-destabilization or tubulin aggregation [34,35]. It is intriguing that the actions of parkin and LRRK2 converge on MTs acetylation and axonal transport, two molecular events that are strictly correlated, since tubulin acetylation speeds up anterograde transport of synaptic vesicles [36] and compensates for the transport defects observed in Huntington's disease [37]. PD-associated Roc-COR domain LRRK2 mutants preferentially bind to deacetylated MTs inhibiting axonal transport in primary neurons of *Drosophila melanogaster* and causing locomotor deficits; interestingly, both the genetic and pharmacological increasing of MT acetylation is able to rescue these defects *in vivo* [38]. Our unpublished results suggest that parkin knockout mouse shows an accelerated accumulation of acetylated MTs, which is specifically associated to dopaminergic neurons, and which can easily lead to the block of mitochondrial transport we observed [Cartelli et al., unpublished]. We have now to remember that both neuronal health [2] and axonal transport [36,37] are sensitive to either overly dynamic (deacetylated) or hyper-stable (heavily acetylated) MTs, as the ones that overcome the lower or the upper MT stability-limiting physiological threshold. Therefore, all these results reinforce the idea that the fine tuning of MT stability, and in particular the modulation of MT acetylation inside dopaminergic neurons, is a critical event during brain aging, and suggest that its failure can be the crossroad between normal neuronal aging and neurodegenerative processes.

The above described MT-based neuronal defects can be counteracted by the administration of MT-stabilizing agents, as already suggested [19,39] and demonstrated in many of the studies above reported [26,28,34]. A reliable alternative can be the fine modulation of MT acetylation, which seems to be tightly associated to axonal transport and locomotor defects [16,36,38]. Since the extent of MT acetylation depends on the balance of the antagonistic acetylases and deacetylases activities, its modulation can be achieved by the use of specific stimulators or inhibitors of α TAT1 or Histone Deacetylase 6 (HDAC6) [14]. The latter specifically acts on tubulin [40], and its inhibition exerts neuroprotection in many age-related neurodegenerative disorders [41] and rescues the transport defects observed in Huntington's disease [37]. In agreement, good results have been achieved in PD experimental models by using inhibitors of MT-deacetylases, as reported by Godena et al., [38] or by Outiero and colleagues [42], which can counteract the excessive accumulation of acetylated MTs or, in other words, the unusual MT hyper-stabilization. Since acetylation marks old MTs and older MTs are more prone to depolymerise than younger ones [12], a further possibility could be trying to induce a sort of MT rejuvenation. This aim does not strictly require the inhibition of MT acetylation but, in order to reach this goal, one could simply stimulate tubulin PTMs associated to newly synthesized and highly dynamic

MTs, as a tubulin tyrosination, which should have beneficial effects also on the neuron itself. Indeed, the specific and unique Tubulin Tyrosine Ligase (TTL) enzyme increases the levels of tyrosinated tubulin at the axonal injury site, activating pro-regenerative transcriptional program important for axon regrowth [43].

Putting all the pieces together, here we propose that the regulation of MT stability, and in particular the modulation of MT acetylation, can be the Achilles' heel of the dopaminergic system, as its weakness could actually discriminate between aging and degeneration. A remaining question is how the disruption of such a ubiquitous process can affect selected neuronal populations and induce specific neurological phenotypes. Our personal feeling is that the answer has to be looked for in the anatomy of neuronal subtypes; indeed, dopaminergic neurons have very long axons whose maintenance and function depend on the correct executions of axonal transport processes and, thus, on MT health, whatever it does mean. In agreement, other long-distance projecting neuronal populations, as striatal and motor neurons, do show alterations of MT stability regulation in pathological contexts [37,44,45]. Therefore, we firmly believe that the failure of a crucial and fine regulated process during brain aging, like the control of MT stability is supposed to be, can specifically affect those neuronal subpopulations whose requirements for MT integrity is pressing and, therefore, it could induce pathological states.

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